

1	5	10
acc gcc gct tct ggt cct ttg gtg gac tac cta tgg atg ctc atc ctg 161		
Thr Ala Ala Ser Gly Pro Leu Val Asp Tyr Leu Trp Met Leu Ile Leu		
15	20	25
ggc ttc att att gca ttt gtc ttg gca ttc tcc gtg gga gcc aat gat 209		
Gly Phe Ile Ile Ala Phe Val Leu Ala Phe Ser Val Gly Ala Asn Asp		
30	35	40
gta gca aat tct ttt ggt aca gct gtg ggc tca ggt gta gtg acc ctg 257		
Val Ala Asn Ser Phe Gly Thr Ala Val Gly Ser Gly Val Val Thr Leu		
45	50	55
aag caa gcc tgc atc cta gct agc atc ttt gaa aca gtc ggc tct gtc 305		
Lys Gln Ala Cys Ile Leu Ala Ser Ile Phe Glu Thr Val Gly Ser Val		
60	65	70
60 70 75		
tta ctg ggg gcc aaa gtc agc gaa acc atc cgg aag ggc ttg att gac 353		
Leu Leu Gly Ala Lys Val Ser Glu Thr Ile Arg Lys Gly Leu Ile Asp		
80	85	90
gtg gag atg tac aac tcg act caa ggg ctg ctg atg gcc ggc tca gtc 401		
Val Glu Met Tyr Asn Ser Thr Gln Gly Leu Leu Met Ala Gly Ser Val		
95	100	105
95 100 105		
agt gct atg ttt ggt tct gct gtg tgg caa ctc gtg gct tcg ttt ttg 449		
Ser Ala Met Phe Gly Ser Ala Val Trp Gln Leu Val Ala Ser Phe Leu		
110	115	120
110 115 120		

aag ctc cct att tct gga acc cat tgt att gtt ggt gca act att ggt 497
 Lys Leu Pro Ile Ser Gly Thr His Cys Ile Val Gly Ala Thr Ile Gly
 125 130 135

ttc tcc ctc gtg gca aag ggg cag gag ggt gtc aag tgg tct gaa ctg 545
 Phe Ser Leu Val Ala Lys Gly Gln Glu Gly Val Lys Trp Ser Glu Leu
 140 145 150 155

ata aaa att gtg atg tct tgg ttc gtg tcc cca ctg ctt tct gga att 593
 Ile Lys Ile Val Met Ser Trp Phe Val Ser Pro Leu Leu Ser Gly Ile
 160 165 170

atg tct gga att tta ttc ttc ctg gtt cgt gca ttc atc ctc cat aag 641
 Met Ser Gly Ile Leu Phe Phe Leu Val Arg Ala Phe Ile Leu His Lys
 175 180 185

gca gat cca gtt cct aat ggt ttg cga gct ttg cca gtt ttc tat gcc 689
 Ala Asp Pro Val Pro Asn Gly Leu Arg Ala Leu Pro Val Phe Tyr Ala
 190 195 200

tgc aca gtt gga ata aac ctc ttt tcc atc atg tat act gga gca ccg 737
 Cys Thr Val Gly Ile Asn Leu Phe Ser Ile Met Tyr Thr Gly Ala Pro
 205 210 215

ttg ctg ggc ttt gac aaa ctt cct ctg tgg ggt acc atc ctc atc tcg 785
 Leu Leu Gly Phe Asp Lys Leu Pro Leu Trp Gly Thr Ile Leu Ile Ser
 220 225 230 235

gtg gga tgt gca gtt ttc tgt gcc ctt atc gtc tgg ttc ttt gta tgt 833
 Val Gly Cys Ala Val Phe Cys Ala Leu Ile Val Trp Phe Phe Val Cys
 240 245 250

ccc agg atg aag aga aaa att gaa cga gaa ata aag tgt agt cct tct 881
 Pro Arg Met Lys Arg Lys Ile Glu Arg Glu Ile Lys Cys Ser Pro Ser
 255 260 265

gaa agc ccc tta atg gaa aaa aag aat agc ttg aaa gaa gac cat gaa 929
 Glu Ser Pro Leu Met Glu Lys Lys Asn Ser Leu Lys Glu Asp His Glu
 270 275 280

gaa aca aag ttg tct gtt ggt gat att gaa aac aag cat cct gtt tct 977
 Glu Thr Lys Leu Ser Val Gly Asp Ile Glu Asn Lys His Pro Val Ser
 285 290 295

gag gta ggg cct gcc act gtg ccc ctc cag gct gtg gtg gag gag aga 1025
 Glu Val Gly Pro Ala Thr Val Pro Leu Gln Ala Val Val Glu Glu Arg
 300 305 310 315

aca gtc tca ttc aaa ctt gga gat ttg gag gaa gct cca gag aga gag 1073
 Thr Val Ser Phe Lys Leu Gly Asp Leu Glu Glu Ala Pro Glu Arg Glu
 320 325 330

agg ctt ccc agc gtg gac ttg aaa gag gaa acc acc agc ata gat agc acc 1121
 Arg Leu Pro Ser Val Asp Leu Lys Glu Glu Thr Ser Ile Asp Ser Thr
 335 340 345

gtg aat ggt gca gtg cag ttg cct aat ggg aac ctt gtc cag ttc agt 1169

Val Asn Gly Ala Val Gln Leu Pro Asn Gly Asn Leu Val Gln Phe Ser

350

355

360

caa gcc gtc agc aac caa ata aac tcc agt ggc cac tac cag tat cac 1217

Gln Ala Val Ser Asn Gln Ile Asn Ser Ser Gly His Tyr Gln Tyr His

365

370

375

acc gtg cat aag gat tcc ggc ctg tac aaa gag cta ctc cat aaa tta 1265

Thr Val His Lys Asp Ser Gly Leu Tyr Lys Glu Leu Leu His Lys Leu

380

385

390

395

cat ctt gcc aag gtg gga gat tgc atg gga gac tcc ggt gac aaa ccc 1313

His Leu Ala Lys Val Gly Asp Cys Met Gly Asp Ser Gly Asp Lys Pro

400

405

410

tta agg cgc aat aat agc tat act tcc tat acc atg gca ata tgt ggc 1361

Leu Arg Arg Asn Asn Ser Tyr Thr Ser Tyr Thr Met Ala Ile Cys Gly

415

420

425

atg cct ctg gat tca ttc cgt gcc aaa gaa ggt gaa cag aag ggc gaa 1409

Met Pro Leu Asp Ser Phe Arg Ala Lys Glu Gly Glu Gln Lys Gly Glu

430

435

440

gaa atg gag aag ctg aca tgg cct aat gca gac tcc aag aag cga att 1457

Glu Met Glu Lys Leu Thr Trp Pro Asn Ala Asp Ser Lys Lys Arg Ile

445

450

455

cga atg gac agt tac acc agt tac tgc aat gct gtg tct gac ctt cac 1505

Arg Met Asp Ser Tyr Thr Ser Tyr Cys Asn Ala Val Ser Asp Leu His

460

465

470

475

tca gca tct gag ata gac atg agt gtc aag gca gag atg ggt cta ggt 1553
 Ser Ala Ser Glu Ile Asp Met Ser Val Lys Ala Glu Met Gly Leu Gly

480

485

490

gac aga aaa gga agt aat ggc tct cta gaa gaa tgg tat gac cag gat 1601
 Asp Arg Lys Gly Ser Asn Gly Ser Leu Glu Glu Trp Tyr Asp Gln Asp

495

500

505

aag cct gaa gtc tct ctc ctc ttc cag ttc ctg cag atc ctt aca gcc 1649
 Lys Pro Glu Val Ser Leu Leu Phe Gln Phe Leu Gln Ile Leu Thr Ala

510

515

520

tgc ttt ggg tca ttc gcc cat ggt ggc aat gac gta agc aat gcc att 1697
 Cys Phe Gly Ser Phe Ala His Gly Gly Asn Asp Val Ser Asn Ala Ile

525

530

535

ggg cct ctg gtt gct tta tat ttg gtt tat gac aca gga gat gtt tct 1745
 Gly Pro Leu Val Ala Leu Tyr Leu Val Tyr Asp Thr Gly Asp Val Ser

540

545

550

555

tca aaa gtg gca aca cca ata tgg ctt cta ctc tat ggt ggt gtt ggt 1793
 Ser Lys Val Ala Thr Pro Ile Trp Leu Leu Tyr Gly Gly Val Gly

560

565

570

atc tgt gtt ggt ctg tgg gtt tgg gga aga aga gtt atc cag acc atg 1841
 Ile Cys Val Gly Leu Trp Val Trp Gly Arg Arg Val Ile Gln Thr Met

575

580

585

ggg aag gat ctg aca ccg atc aca ccc tct agt ggc ttc agt att gaa 1889

Gly Lys Asp Leu Thr Pro Ile Thr Pro Ser Ser Gly Phe Ser Ile Glu

590

595

600

ctg gca tct gcc ctc act gtg gtg att gca tca aat att ggc ctt ccc 1937

Leu Ala Ser Ala Leu Thr Val Val Ile Ala Ser Asn Ile Gly Leu Pro

605

610

615

atc agt aca aca cat tgt aaa gtg ggc tct gtt gtg tct gtt ggc tgg 1985

Ile Ser Thr Thr His Cys Lys Val Gly Ser Val Val Ser Val Gly Trp

620

625

630

635

ctc cgg tcc aag aag gct gtt gac tgg cgt ctc ttt cgt aac att ttt 2033

Leu Arg Ser Lys Lys Ala Val Asp Trp Arg Leu Phe Arg Asn Ile Phe

640

645

650

atg gcc tgg ttt gtc aca gtc cct att tct gga gtt atc agt gct gcc 2081

Met Ala Trp Phe Val Thr Val Pro Ile Ser Gly Val Ile Ser Ala Ala

655

660

665

atc atg gca atc ttc aga tat gtc atc ctc aga atg tgaagctgtt 2127

Ile Met Ala Ile Phe Arg Tyr Val Ile Leu Arg Met

670

675

ttagattaaa atttgtgtca atgtttggga ccatcttagg tattcctgct cccctgaaga 2187

atgattacag tgttaacaga agactgacaa gagtctttt atttgggagc cagaggaggg 2247

aagtgttact tgtgctataa ctgctttgt gctaaatatg aattgtctca aaatttagctg 2307

tgtaaaatag cccgggttcc actggctcct gctgaggccc ctttccttc tgggctgtga 2367

atccctgtac atatttctct accttttgta tcaggctca attccattat gtttaatgt 2427

tgtctctgaa gatgacttgt gatTTTTT TCTTTTTT aaaccatgaa gagccgttg 2487

acagagcatg ctctgcgttg ttggTTcac cagcttctgc cctcacatgc acaggattt 2547

aacaacaaaa atataactac aactccctt gtagtctctt atataagtag agtccttgg 2607

actctgccct cctgtcagta gtggcaggat ctattggcat attcgggagc ttcttagagg 2667

gatgaggttc tttgaacaca gtgaaaattt aaatttagtaa ctttttgca agcagtttat 2727

tgactgttat tgctaagaag aagtaagaaa gaaaaagcct gttggcaatc ttggttattt 2787

cttaagatt tctggcagtg tggatggat gaatgaagtg gaatgtgaac tttggcaag 2847

ttaaatggga cagccttcca ttttcatttg tctaccttt aactgaataa aaaagcctac 2907

agtTTTTAG 2916

<210> 125

<211> 288

<212> PRT

<213> Homo sapiens

<400> 125

Met Glu Arg Pro Gln Pro Asp Ser Met Pro Gln Asp Leu Ser Glu Ala

1

5

10

15

Leu Lys Glu Ala Thr Lys Glu Val His Thr Gln Ala Glu Asn Ala Glu

20

25

30

Phe Met Arg Asn Phe Gln Lys Gly Gln Val Thr Arg Asp Gly Phe Lys

35

40

45

Leu Val Met Ala Ser Leu Tyr His Ile Tyr Val Ala Leu Glu Glu Glu

50

55

60

Ile Glu Arg Asn Lys Glu Ser Pro Val Phe Ala Pro Val Tyr Phe Pro

65

70

75

80

Glu Glu Leu His Arg Lys Ala Ala Leu Glu Gln Asp Leu Ala Phe Trp

85

90

95

Tyr Gly Pro Arg Trp Gln Glu Val Ile Pro Tyr Thr Pro Ala Met Gln

100

105

110

Arg Tyr Val Lys Arg Leu His Glu Val Gly Arg Thr Glu Pro Glu Leu

115

120

125

Leu Val Ala His Ala Tyr Thr Arg Tyr Leu Gly Asp Leu Ser Gly Gly

130

135

140

Gln Val Leu Lys Lys Ile Ala Gln Lys Ala Leu Asp Leu Pro Ser Ser
 145 150 155 160

Gly Glu Gly Leu Ala Phe Phe Thr Phe Pro Asn Ile Ala Ser Ala Thr
 165 170 175

Lys Phe Lys Gln Leu Tyr Arg Ser Arg Met Asn Ser Leu Glu Met Thr
 180 185 190

Pro Ala Val Arg Gln Arg Val Ile Glu Glu Ala Lys Thr Ala Phe Leu
 195 200 205

Leu Asn Ile Gln Leu Phe Glu Glu Leu Gln Glu Leu Leu Thr His Asp
 210 215 220

Thr Lys Asp Gln Ser Pro Ser Arg Ala Pro Gly Leu Arg Gln Arg Ala
 225 230 235 240

Ser Asn Lys Val Gln Asp Ser Ala Pro Val Glu Thr Pro Arg Gly Lys
 245 250 255

Pro Pro Leu Asn Thr Arg Ser Gln Ala Pro Leu Leu Arg Trp Val Leu
 260 265 270

Thr Leu Ser Phe Leu Val Ala Thr Val Ala Val Gly Leu Tyr Ala Met
 275 280 285

<210> 126

<211> 1550

<212> DNA

<213> Homo sapiens

<220>

<221> CDS

<222> (81)..(944)

<400> 126

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acgagccag caccggccgg atg gag cgt ccg caa ccc gac agc atg ccc cag 113

Met Glu Arg Pro Gln Pro Asp Ser Met Pro Gln

1

5

10

gat ttg tca gag gcc ctg aag gag gcc acc aag gag gtg cac acc cag 161

Asp Leu Ser Glu Ala Leu Lys Glu Ala Thr Lys Glu Val His Thr Gln

15

20

25

gca gag aat gct gag ttc atg agg aac ttt cag aag ggc cag gtg acc 209

Ala Glu Asn Ala Glu Phe Met Arg Asn Phe Gln Lys Gly Gln Val Thr

30

35

40

cga gac ggc ttc aag ctg gtg atg gcc tcc ctg tac cac atc tat gtg 257

Arg Asp Gly Phe Lys Leu Val Met Ala Ser Leu Tyr His Ile Tyr Val

45

50

55

gcc ctg gag gag att gag cgc aac aag gag agc cca gtc ttc gcc 305

Ala Leu Glu Glu Glu Ile Glu Arg Asn Lys Glu Ser Pro Val Phe Ala

60

65

70

75

cct gtc tac ttc cca gaa gag ctg cac cgc aag gct gcc ctg gag cag 353

Pro Val Tyr Phe Pro Glu Glu Leu His Arg Lys Ala Ala Leu Glu Gln

80

85

90

gac ctg gcc ttc tgg tac ggg ccc cgc tgg cag gag gtc atc ccc tac 401

Asp Leu Ala Phe Trp Tyr Gly Pro Arg Trp Gln Glu Val Ile Pro Tyr

95

100

105

aca cca gcc atg cag cgc tat gtg aag cgg ctc cac gag gtg ggg cgc 449

Thr Pro Ala Met Gln Arg Tyr Val Lys Arg Leu His Glu Val Gly Arg

110

115

120

aca gag ccc gag ctg ctg gtg gcc cac gcc tac acc cgc tac ctg ggt 497

Thr Glu Pro Glu Leu Leu Val Ala His Ala Tyr Thr Arg Tyr Leu Gly

125

130

135

gac ctg tct ggg ggc cag gtg ctc aaa aag att gcc cag aaa gcc ctg 545

Asp Leu Ser Gly Gly Gln Val Leu Lys Lys Ile Ala Gln Lys Ala Leu

140

145

150

155

gac ctg ccc agc tct ggc gag ggc ctg gcc ttc ttc acc ttc ccc aac 593

Asp Leu Pro Ser Ser Gly Glu Gly Leu Ala Phe Phe Thr Phe Pro Asn

160

165

170

att gcc agt gcc acc aag ttc aag cag ctc tac cgc tcc cgc atg aac 641

Ile Ala Ser Ala Thr Lys Phe Lys Gln Leu Tyr Arg Ser Arg Met Asn

175

180

185

tcc ctg gag atg act ccc gca gtc agg cag agg gtg ata gaa gag gcc 689

Ser Leu Glu Met Thr Pro Ala Val Arg Gln Arg Val Ile Glu Glu Ala

190

195

200

aag act gcg ttc ctg ctc aac atc cag ctc ttt gag gag ttg cag gag 737

Lys Thr Ala Phe Leu Leu Asn Ile Gln Leu Phe Glu Glu Leu Gln Glu

205

210

215

ctg ctg acc cat gac acc aag gac cag agc ccc tca cgg gca cca ggg 785

Leu Leu Thr His Asp Thr Lys Asp Gln Ser Pro Ser Arg Ala Pro Gly

220

225

230

235

ctt cgc cag cgg gcc agc aac aaa gtg caa gat tct gcc ccc gtg gag 833

Leu Arg Gln Arg Ala Ser Asn Lys Val Gln Asp Ser Ala Pro Val Glu

240

245

250

act ccc aga ggg aag ccc cca ctc aac acc cgc tcc cag gct ccg ctt 881

Thr Pro Arg Gly Lys Pro Pro Leu Asn Thr Arg Ser Gln Ala Pro Leu

255

260

265

ctc cga tgg gtc ctt aca ctc agc ttt ctg gtg gcg aca gtt gct gta 929

Leu Arg Trp Val Leu Thr Leu Ser Phe Leu Val Ala Thr Val Ala Val

270

275

280

ggg ctt tat gcc atg tgaatgcagg catgctggct cccagggcca tgaaccttgt 984

Gly Leu Tyr Ala Met

285

ccgttggaaag gccttcattc tagagaggga attcttttgg ctggcttcct taccgtgggc 1044

actgaaggct ttcatggcct ccagccctct cactgtgtcc ctctctctgg aaaggaggaa 1104

ggagcctatg gcatcttccc caacgaaaag cacatccagg caatggccta aacttcagag 1164

ggggcgaagg ggtcagccct gcccttcagc atcctcagtt cctgcagcag agcctggaag 1224

acacccta at gtggcagctg tctcaaacct ccaaaagccc tgagttcaa gtatccttgt 1284

tgacacggcc atgaccactt tccccgtggg ccatggcaat ttttacacaa acctgaaaag 1344

atgttgtgtc ttgtgtttt gtcttatttt tggtggagcc actctgttcc tggctcagcc 1404

tcaaatgcag tattttgtt gtgttctgtt gttttatag cagggttggg gtggttttt 1464

agccatgcgt gggtggggag ggaggtgttt aacggcactg tggccttgg ctaactttt 1524

tgtgaaataa taaacaacat tgtctg 1550

<210> 127

<211> 135

<212> PRT

<213> Homo sapiens

<400> 127

Met Ala Cys Gly Leu Val Ala Ser Asn Leu Asn Leu Lys Pro Gly Glu

1

5

10

15

Cys Leu Arg Val Arg Gly Glu Val Ala Pro Asp Ala Lys Ser Phe Val

20

25

30

Leu Asn Leu Gly Lys Asp Ser Asn Asn Leu Cys Leu His Phe Asn Pro

35

40

45

Arg Phe Asn Ala His Gly Asp Ala Asn Thr Ile Val Cys Asn Ser Lys

50

55

60

Asp Gly Gly Ala Trp Gly Thr Glu Gln Arg Glu Ala Val Phe Pro Phe

65

70

75

80

Gln Pro Gly Ser Val Ala Glu Val Cys Ile Thr Phe Asp Gln Ala Asn

85

90

95

Leu Thr Val Lys Leu Pro Asp Gly Tyr Glu Phe Lys Phe Pro Asn Arg

100

105

110

Leu Asn Leu Glu Ala Ile Asn Tyr Met Ala Ala Asp Gly Asp Phe Lys

115

120

125

Ile Lys Cys Val Ala Phe Asp

130

135

<210> 128

<211> 507

<212> DNA

<213> Homo sapiens

<220>

<221> CDS

<222> (50)..(454)

<400> 128

cttctgacag ctggcgcc tgccggaa catcctcctg gactcaatc atg gct tgt 58

Met Ala Cys

1

ggc ctc gtc gcc agc aac ctt aat ctc aaa cct gga gag tgc ctt cga 106

Gly Leu Val Ala Ser Asn Leu Asn Leu Lys Pro Gly Glu Cys Leu Arg

5

10

15

gtg cga ggc gag gtg gct cct gac gct aag agc ttc gtg ctg aac ctg 154

Val Arg Gly Glu Val Ala Pro Asp Ala Lys Ser Phe Val Leu Asn Leu

20

25

30

35

ggc aaa gac agc aac aac ctg tgc ctg cac ttc aac cct cgc ttc aac 202

Gly Lys Asp Ser Asn Asn Leu Cys Leu His Phe Asn Pro Arg Phe Asn

40

45

50

gcc cac ggc gac gcc aac acc atc gtg tgc aac agc aag gac ggc ggg 250

Ala His Gly Asp Ala Asn Thr Ile Val Cys Asn Ser Lys Asp Gly Gly

55

60

65

gcc tgg ggg acc gag cag cgg gag gct gtc ttt ccc ttc cag cct gga 298

Ala Trp Gly Thr Glu Gln Arg Glu Ala Val Phe Pro Phe Gln Pro Gly

70

75

80

agt gtt gca gag gtg tgc atc acc ttc gac cag gcc aac ctg acc gtc 346

Ser Val Ala Glu Val Cys Ile Thr Phe Asp Gln Ala Asn Leu Thr Val

85

90

95

aag ctg cca gat gga tac gaa ttc aag ttc ccc aac cgc ctc aac ctg 394

Lys Leu Pro Asp Gly Tyr Glu Phe Lys Phe Pro Asn Arg Leu Asn Leu

100

105

110

115

gag gcc atc aac tac atg gca gct gac ggt gac ttc aag atc aaa tgt 442

Glu Ala Ile Asn Tyr Met Ala Ala Asp Gly Asp Phe Lys Ile Lys Cys

120

125

130

gtg gcc ttt gac tgaaatcagc cagcccatgg cccccataa aggcagctgc 494

Val Ala Phe Asp

135

ctctgctccc ctg

507

<210> 129

<211> 662

<212> PRT

<213> Homo sapiens

<400> 129

Met Asn Lys Glu Ile Pro Asn Gly Asn Thr Ser Glu Leu Ile Phe Asn

1

5

10

15

Ala Val His Val Lys Asp Ala Gly Phe Tyr Val Cys Arg Val Asn Asn

20

25

30

Asn Phe Thr Phe Glu Phe Ser Gln Trp Ser Gln Leu Asp Val Cys Asp

35

40

45

Ile Pro Glu Ser Phe Gln Arg Ser Val Asp Gly Val Ser Glu Ser Lys

50

55

60

Leu Gln Ile Cys Val Glu Pro Thr Ser Gln Lys Leu Met Pro Gly Ser

65

70

75

80

Thr Leu Val Leu Gln Cys Val Ala Val Gly Ser Pro Ile Pro His Tyr

85

90

95

Gln Trp Phe Lys Asn Glu Leu Pro Leu Thr His Glu Thr Lys Lys Leu

100

105

110

Tyr Met Val Pro Tyr Val Asp Leu Glu His Gln Gly Thr Tyr Trp Cys

115

120

125

His Val Tyr Asn Asp Arg Asp Ser Gln Asp Ser Lys Lys Val Glu Ile

130

135

140

Ile Ile Gly Arg Thr Asp Glu Ala Val Glu Cys Thr Glu Asp Glu Leu

145 150 155 160

Asn Asn Leu Gly His Pro Asp Asn Lys Glu Gln Thr Thr Asp Gln Pro

165	170	175
Leu Ala Lys Asp Lys Val Ala Leu Leu Ile Gly Asn Met Asn Tyr Arg		
180	185	190
Glu His Pro Lys Leu Lys Ala Pro Leu Val Asp Val Tyr Glu Leu Thr		
195	200	205
Asn Leu Leu Arg Gln Leu Asp Phe Lys Val Val Ser Leu Leu Asp Leu		
210	215	220
Thr Glu Tyr Glu Met Arg Asn Ala Val Asp Glu Phe Leu Leu Leu Leu		
225	230	235
Asp Lys Gly Val Tyr Gly Leu Leu Tyr Tyr Ala Gly His Gly Tyr Glu		
245	250	255
Asn Phe Gly Asn Ser Phe Met Val Pro Val Asp Ala Pro Asn Pro Tyr		
260	265	270
Arg Ser Glu Asn Cys Leu Cys Val Gln Asn Ile Leu Lys Leu Met Gln		
275	280	285
Glu Lys Glu Thr Gly Leu Asn Val Phe Leu Leu Asp Met Cys Arg Lys		
290	295	300
Arg Asn Asp Tyr Asp Asp Thr Ile Pro Ile Leu Asp Ala Leu Lys Val		
305	310	315
320		

Thr Ala Asn Ile Val Phe Gly Tyr Ala Thr Cys Gln Gly Ala Glu Ala

325

330

335

Phe Glu Ile Gln His Ser Gly Leu Ala Asn Gly Ile Phe Met Lys Phe

340

345

350

Leu Lys Asp Arg Leu Leu Glu Asp Lys Lys Ile Thr Val Leu Leu Asp

355

360

365

Glu Val Ala Glu Asp Met Gly Lys Cys His Leu Thr Lys Gly Lys Gln

370

375

380

Ala Leu Glu Ile Arg Ser Ser Leu Ser Glu Lys Arg Ala Leu Thr Asp

385

390

395

400

Pro Ile Gln Gly Thr Glu Tyr Ser Ala Glu Ser Leu Val Arg Asn Leu

405

410

415

Gln Trp Ala Lys Ala His Glu Leu Pro Glu Ser Met Cys Leu Lys Phe

420

425

430

Asp Cys Gly Val Gln Ile Gln Leu Gly Phe Ala Ala Glu Phe Ser Asn

435

440

445

Val Met Ile Ile Tyr Thr Ser Ile Val Tyr Lys Pro Pro Glu Ile Ile

450

455

460

Met Cys Asp Ala Tyr Val Thr Asp Phe Pro Leu Asp Leu Asp Ile Asp

465

470

475

480

Pro Lys Asp Ala Asn Lys Gly Thr Pro Glu Glu Thr Gly Ser Tyr Leu

485

490

495

Val Ser Lys Asp Leu Pro Lys His Cys Leu Tyr Thr Arg Leu Ser Ser

500

505

510

Leu Gln Lys Leu Lys Glu His Leu Val Phe Thr Val Cys Leu Ser Tyr

515

520

525

Gln Tyr Ser Gly Leu Glu Asp Thr Val Glu Asp Lys Gln Glu Val Asn

530

535

540

Val Gly Lys Pro Leu Ile Ala Lys Leu Asp Met His Arg Gly Leu Gly

545

550

555

560

Arg Lys Thr Cys Phe Gln Thr Cys Leu Met Ser Asn Gly Pro Tyr Gln

565

570

575

Ser Ser Ala Ala Thr Ser Gly Gly Ala Gly His Tyr His Ser Leu Gln

580

585

590

Asp Pro Phe His Gly Val Tyr His Ser His Pro Gly Asn Pro Ser Asn

595

600

605

Val Thr Pro Ala Asp Ser Cys His Cys Ser Arg Thr Pro Asp Ala Phe

610

615

620

Ile Ser Ser Phe Ala His His Ala Ser Cys His Phe Ser Arg Ser Asn

625

630

635

640

Val Pro Val Glu Thr Thr Asp Glu Ile Pro Phe Ser Phe Ser Asp Arg

645

650

655

Leu Arg Ile Ser Glu Lys

660

<210> 130

<211> 2251

<212> DNA

<213> Homo sapiens

<220>

<221> CDS

<222> (74)..(2059)

<400> 130

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tcagtggttc aaa atg aat aaa gag att cca aat gga aat aca tca gag 109

Met Asn Lys Glu Ile Pro Asn Gly Asn Thr Ser Glu

1

5

10

ctt att ttt aat gca gtg cat gta aaa gat gca ggc ttt tat gtc tgt 157

Leu Ile Phe Asn Ala Val His Val Lys Asp Ala Gly Phe Tyr Val Cys

15

20

25

cga gtt aat aac aat ttc acc ttt gaa ttc agc cag tgg tca cag ctg 205
 Arg Val Asn Asn Asn Phe Thr Phe Glu Phe Ser Gln Trp Ser Gln Leu
 30 35 40

gat gtt tgc gac atc cca gag agc ttc cag aga agt gtt gat ggc gtc 253
 Asp Val Cys Asp Ile Pro Glu Ser Phe Gln Arg Ser Val Asp Gly Val
 45 50 55 60

tct gaa tcc aag ttg caa atc tgt gtt gaa cca act tcc caa aag ctg 301
 Ser Glu Ser Lys Leu Gln Ile Cys Val Glu Pro Thr Ser Gln Lys Leu
 65 70 75

atg cca ggc agc aca ttg gtt tta cag tgt gtt gct gtt gga agc cct 349
 Met Pro Gly Ser Thr Leu Val Leu Gln Cys Val Ala Val Gly Ser Pro
 80 85 90

att cct cac tac cag tgg ttc aaa aat gaa tta cca tta aca cat gag 397
 Ile Pro His Tyr Gln Trp Phe Lys Asn Glu Leu Pro Leu Thr His Glu
 95 100 105

acc aaa aag cta tac atg gtg cct tat gtg gat ttg gaa cac caa gga 445
 Thr Lys Lys Leu Tyr Met Val Pro Tyr Val Asp Leu Glu His Gln Gly
 110 115 120

acc tac tgg tgt cat gta tat aat gat cga gac agt caa gat agc aag 493
 Thr Tyr Trp Cys His Val Tyr Asn Asp Arg Asp Ser Gln Asp Ser Lys
 125 130 135 140

aag gta gaa atc atc ata gga aga aca gat gag gca gtg gag tgc act 541

Lys Val Glu Ile Ile Gly Arg Thr Asp Glu Ala Val Glu Cys Thr

145

150

155

gaa gat gaa tta aat aat ctt ggt cat cct gat aat aaa gag caa aca 589

Glu Asp Glu Leu Asn Asn Leu Gly His Pro Asp Asn Lys Glu Gln Thr

160

165

170

act gac cag cct ttg gcg aag gac aag gtt gcc ctt ttg ata gga aat 637

Thr Asp Gln Pro Leu Ala Lys Asp Lys Val Ala Leu Leu Ile Gly Asn

175

180

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atg aat tac cgg gag cac ccc aag ctc aaa gct cct ttg gtg gat gtg 685

Met Asn Tyr Arg Glu His Pro Lys Leu Lys Ala Pro Leu Val Asp Val

190

195

200

tac gaa ttg act aac tta ctg aga cag ctg gac ttc aaa gtg gtt tca 733

Tyr Glu Leu Thr Asn Leu Leu Arg Gln Leu Asp Phe Lys Val Val Ser

205

210

215

220

ctg ttg gat ctt act gaa tat gag atg cgt aat gct gtg gat gag ttt 781

Leu Leu Asp Leu Thr Glu Tyr Glu Met Arg Asn Ala Val Asp Glu Phe

225

230

235

tta ctc ctt tta gac aag gga gta tat ggg tta tta tat tat gca gga 829

Leu Leu Leu Asp Lys Gly Val Tyr Gly Leu Leu Tyr Tyr Ala Gly

240

245

250

cat ggt tat gaa aat ttt ggg aac agc ttc atg gtc ccc gtt gat gct 877

His Gly Tyr Glu Asn Phe Gly Asn Ser Phe Met Val Pro Val Asp Ala

255

260

265

cca aat cca tat agg tct gaa aat tgt ctg tgt gta caa aat ata ctg 925

Pro Asn Pro Tyr Arg Ser Glu Asn Cys Leu Cys Val Gln Asn Ile Leu

270

275

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aaa ttg atg caa gaa aaa gaa act gga ctt aat gtg ttc tta ttg gat 973

Lys Leu Met Gln Glu Lys Glu Thr Gly Leu Asn Val Phe Leu Leu Asp

285

290

295

300

atg tgt agg aaa aga aat gac tac gat gat acc att cca atc ttg gat 1021

Met Cys Arg Lys Arg Asn Asp Tyr Asp Asp Thr Ile Pro Ile Leu Asp

305

310

315

gca cta aaa gtc acc gcc aat att gtg ttt gga tat gcc acg tgt caa 1069

Ala Leu Lys Val Thr Ala Asn Ile Val Phe Gly Tyr Ala Thr Cys Gln

320

325

330

gga gca gaa gct ttt gaa atc cag cat tct gga ttg gca aat gga atc 1117

Gly Ala Glu Ala Phe Glu Ile Gln His Ser Gly Leu Ala Asn Gly Ile

335

340

345

ttt atg aaa ttt tta aaa gac aga tta tta gaa gat aag aaa atc act 1165

Phe Met Lys Phe Leu Lys Asp Arg Leu Leu Glu Asp Lys Lys Ile Thr

350

355

360

gtg tta ctg gat gaa gtt gca gaa gat atg ggt aag tgt cac ctt acc 1213

Val Leu Leu Asp Glu Val Ala Glu Asp Met Gly Lys Cys His Leu Thr

365

370

375

380

aaa ggc aaa cag gct cta gag att cga agt agt tta tct gag aag aga 1261
 Lys Gly Lys Gln Ala Leu Glu Ile Arg Ser Ser Leu Ser Glu Lys Arg

385 390 395

gca ctt act gat cca ata cag gga aca gaa tat tct gct gaa tct ctt 1309
 Ala Leu Thr Asp Pro Ile Gln Gly Thr Glu Tyr Ser Ala Glu Ser Leu

400 405 410

gtg cgg aat cta cag tgg gcc aag gct cat gaa ctt cca gaa agt atg 1357
 Val Arg Asn Leu Gln Trp Ala Lys Ala His Glu Leu Pro Glu Ser Met

415 420 425

tgt ctt aag ttt gac tgt ggt gtt cag att caa tta gga ttt gca gct 1405
 Cys Leu Lys Phe Asp Cys Gly Val Gln Ile Gln Leu Gly Phe Ala Ala
 430 435 440

gag ttt tcc aat gtc atg atc atc tat aca agt ata gtt tac aaa cca 1453
 Glu Phe Ser Asn Val Met Ile Ile Tyr Thr Ser Ile Val Tyr Lys Pro
 445 450 455 460

ccg gag ata ata atg tgt gat gcc tac gtt act gat ttt cca ctt gat 1501
 Pro Glu Ile Ile Met Cys Asp Ala Tyr Val Thr Asp Phe Pro Leu Asp
 465 470 475

cta gat att gat cca aaa gat gca aat aaa ggc aca cct gaa gaa act 1549
 Leu Asp Ile Asp Pro Lys Asp Ala Asn Lys Gly Thr Pro Glu Glu Thr
 480 485 490

ggc agc tac ttg gta tca aag gat ctt ccc aag cat tgc ctc tat acc 1597
 Gly Ser Tyr Leu Val Ser Lys Asp Leu Pro Lys His Cys Leu Tyr Thr
 495 500 505

aga ctc agt tca ctg caa aaa tta aag gaa cat cta gtc ttc aca gta 1645
 Arg Leu Ser Ser Leu Gln Lys Leu Lys Glu His Leu Val Phe Thr Val
 510 515 520

tgt tta tca tat cag tac tca gga ttg gaa gat act gta gag gac aag 1693
 Cys Leu Ser Tyr Gln Tyr Ser Gly Leu Glu Asp Thr Val Glu Asp Lys
 525 530 535 540

cag gaa gtg aat gtt ggg aaa cct ctc att gct aaa tta gac atg cat 1741
 Gln Glu Val Asn Val Gly Lys Pro Leu Ile Ala Lys Leu Asp Met His
 545 550 555

cga ggt ttg gga agg aag act tgc ttt caa act tgt ctt atg tct aat 1789
 Arg Gly Leu Gly Arg Lys Thr Cys Phe Gln Thr Cys Leu Met Ser Asn
 560 565 570

ggt cct tac cag agt tct gca gcc acc tca gga gga gca ggg cat tat 1837
 Gly Pro Tyr Gln Ser Ser Ala Ala Thr Ser Gly Gly Ala Gly His Tyr
 575 580 585

cac tca ttg caa gac cca ttc cat ggt gtt tac cat tca cat cct ggt 1885
 His Ser Leu Gln Asp Pro Phe His Gly Val Tyr His Ser His Pro Gly
 590 595 600

aat cca agt aat gtt aca cca gca gat agc tgt cat tgc agc cgg act 1933

Asn Pro Ser Asn Val Thr Pro Ala Asp Ser Cys His Cys Ser Arg Thr

605

610

615

620

cca gat gca ttt att tca agt ttc gct cac cat gct tca tgt cat ttt 1981

Pro Asp Ala Phe Ile Ser Ser Phe Ala His His Ala Ser Cys His Phe

625

630

635

agt aga agt aat gtg cca gta gag aca act gat gaa ata cca ttt agt 2029

Ser Arg Ser Asn Val Pro Val Glu Thr Thr Asp Glu Ile Pro Phe Ser

640

645

650

ttc tct gac agg ctc aga att tct gaa aaa tgacctcctt gttttgaaa 2079

Phe Ser Asp Arg Leu Arg Ile Ser Glu Lys

655

660

gttagcataa tttagatgc ctgtgaaata gtactgcact tacataaaat gagacattgt 2139

aaaaaggcaa atttgtatat gtagagaaag aatagtagta actgtttcat agcaaacttc 2199

aggacttga gatgttggaa ttacatttt taattacaga cttcctctt ct

2251

<210> 131

<211> 824

<212> PRT

<213> Homo sapiens

<400> 131

Met Ser Leu Leu Gly Asp Pro Leu Gln Ala Leu Pro Pro Ser Ala Ala

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Pro Thr Gly Pro Leu Leu Ala Pro Pro Ala Gly Ala Thr Leu Asn Arg

20

25

30

Leu Arg Glu Pro Leu Leu Arg Arg Leu Ser Glu Leu Leu Asp Gln Ala

35

40

45

Pro Glu Gly Arg Gly Trp Arg Arg Leu Ala Glu Leu Ala Gly Ser Arg

50

55

60

Gly Arg Leu Arg Leu Ser Cys Leu Asp Leu Glu Gln Cys Ser Leu Lys

65

70

75

80

Val Leu Glu Pro Glu Gly Ser Pro Ser Leu Cys Leu Leu Lys Leu Met

85

90

95

Gly Glu Lys Gly Cys Thr Val Thr Glu Leu Ser Asp Phe Leu Gln Ala

100

105

110

Met Glu His Thr Glu Val Leu Gln Leu Leu Ser Pro Pro Gly Ile Lys

115

120

125

Ile Thr Val Asn Pro Glu Ser Lys Ala Val Leu Ala Gly Gln Phe Val

130

135

140

Lys Leu Cys Cys Arg Ala Thr Gly His Pro Phe Val Gln Tyr Gln Trp

145

150

155

160

Phe Lys Met Asn Lys Glu Ile Pro Asn Gly Asn Thr Ser Glu Leu Ile

165 170 175

Phe Asn Ala Val His Val Lys Asp Ala Gly Phe Tyr Val Cys Arg Val

180 185 190

Asn Asn Asn Phe Thr Phe Glu Phe Ser Gln Trp Ser Gln Leu Asp Val

195 200 205

Cys Asp Ile Pro Glu Ser Phe Gln Arg Ser Val Asp Gly Val Ser Glu

210 215 220

Ser Lys Leu Gln Ile Cys Val Glu Pro Thr Ser Gln Lys Leu Met Pro

225 230 235 240

Gly Ser Thr Leu Val Leu Gln Cys Val Ala Val Gly Ser Pro Ile Pro

245 250 255

His Tyr Gln Trp Phe Lys Asn Glu Leu Pro Leu Thr His Glu Thr Lys

260 265 270

Lys Leu Tyr Met Val Pro Tyr Val Asp Leu Glu His Gln Gly Thr Tyr

275 280 285

Trp Cys His Val Tyr Asn Asp Arg Asp Ser Gln Asp Ser Lys Lys Val

290 295 300

Glu Ile Ile Ile Gly Arg Thr Asp Glu Ala Val Glu Cys Thr Glu Asp

305 310 315 320

Glu Leu Asn Asn Leu Gly His Pro Asp Asn Lys Glu Gln Thr Thr Asp

325

330

335

Gln Pro Leu Ala Lys Asp Lys Val Ala Leu Leu Ile Gly Asn Met Asn

340

345

350

Tyr Arg Glu His Pro Lys Leu Lys Ala Pro Leu Val Asp Val Tyr Glu

355

360

365

Leu Thr Asn Leu Leu Arg Gln Leu Asp Phe Lys Val Val Ser Leu Leu

370

375

380

Asp Leu Thr Glu Tyr Glu Met Arg Asn Ala Val Asp Glu Phe Leu Leu

385

390

395

400

Leu Leu Asp Lys Gly Val Tyr Gly Leu Leu Tyr Tyr Ala Gly His Gly

405

410

415

Tyr Glu Asn Phe Gly Asn Ser Phe Met Val Pro Val Asp Ala Pro Asn

420

425

430

Pro Tyr Arg Ser Glu Asn Cys Leu Cys Val Gln Asn Ile Leu Lys Leu

435

440

445

Met Gln Glu Lys Glu Thr Gly Leu Asn Val Phe Leu Leu Asp Met Cys

450

455

460

Arg Lys Arg Asn Asp Tyr Asp Asp Thr Ile Pro Ile Leu Asp Ala Leu

465	470	475	480
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Lys Val Thr Ala Asn Ile Val Phe Gly Tyr Ala Thr Cys Gln Gly Ala

485	490	495
-----	-----	-----

Glu Ala Phe Glu Ile Gln His Ser Gly Leu Ala Asn Gly Ile Phe Met

500	505	510
-----	-----	-----

Lys Phe Leu Lys Asp Arg Leu Leu Glu Asp Lys Lys Ile Thr Val Leu

515	520	525
-----	-----	-----

Leu Asp Glu Val Ala Glu Asp Met Gly Lys Cys His Leu Thr Lys Gly

530	535	540
-----	-----	-----

Lys Gln Ala Leu Glu Ile Arg Ser Ser Leu Ser Glu Lys Arg Ala Leu

545	550	555	560
-----	-----	-----	-----

Thr Asp Pro Ile Gln Gly Thr Glu Tyr Ser Ala Glu Ser Leu Val Arg

565	570	575
-----	-----	-----

Asn Leu Gln Trp Ala Lys Ala His Glu Leu Pro Glu Ser Met Cys Leu

580	585	590
-----	-----	-----

Lys Phe Asp Cys Gly Val Gln Ile Gln Leu Gly Phe Ala Ala Glu Phe

595	600	605
-----	-----	-----

Ser Asn Val Met Ile Ile Tyr Thr Ser Ile Val Tyr Lys Pro Pro Glu

610	615	620
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Ile Ile Met Cys Asp Ala Tyr Val Thr Asp Phe Pro Leu Asp Leu Asp
625 630 635 640

Ile Asp Pro Lys Asp Ala Asn Lys Gly Thr Pro Glu Glu Thr Gly Ser
645 650 655

Tyr Leu Val Ser Lys Asp Leu Pro Lys His Cys Leu Tyr Thr Arg Leu
660 665 670

Ser Ser Leu Gln Lys Leu Lys Glu His Leu Val Phe Thr Val Cys Leu
675 680 685

Ser Tyr Gln Tyr Ser Gly Leu Glu Asp Thr Val Glu Asp Lys Gln Glu
690 695 700

Val Asn Val Gly Lys Pro Leu Ile Ala Lys Leu Asp Met His Arg Gly
705 710 715 720

Leu Gly Arg Lys Thr Cys Phe Gln Thr Cys Leu Met Ser Asn Gly Pro
725 730 735

Tyr Gln Ser Ser Ala Ala Thr Ser Gly Gly Ala Gly His Tyr His Ser
740 745 750

Leu Gln Asp Pro Phe His Gly Val Tyr His Ser His Pro Gly Asn Pro
755 760 765

Ser Asn Val Thr Pro Ala Asp Ser Cys His Cys Ser Arg Thr Pro Asp
770 775 780

Ala Phe Ile Ser Ser Phe Ala His His Ala Ser Cys His Phe Ser Arg

785 790 795 800

Ser Asn Val Pro Val Glu Thr Thr Asp Glu Ile Pro Phe Ser Phe Ser

805 810 815

Asp Arg Leu Arg Ile Ser Glu Lys

820

<210> 132

<211> 2828

<212> DNA

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<220>

<221> CDS

<222> (165)..(2636)

<400> 132

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gccccctgcct ccgcggctcg gaggcgagcg gaaggtgccccc cggggccgag gcccgtgacg 120

gggcgggcgg gagccccggc agtccgggt cgccggcgag ggcc atg tcg ctg ttg 176

Met Ser Leu Leu

ggg gac ccg cta cag gcc ctg ccg ccc tcg gcc gcc ccc acg ggg ccg 224
 Gly Asp Pro Leu Gln Ala Leu Pro Pro Ser Ala Ala Pro Thr Gly Pro
 5 10 15 20

ctg ctc gcc cct ccg gcc ggc gcg acc ctc aac cgc ctg cgg gag ccg 272
 Leu Leu Ala Pro Pro Ala Gly Ala Thr Leu Asn Arg Leu Arg Glu Pro
 25 30 35

ctg ctg cgg agg ctc agc gag ctc ctg gat cag gcg ccc gag ggc cgg 320
 Leu Leu Arg Arg Leu Ser Glu Leu Leu Asp Gln Ala Pro Glu Gly Arg
 40 45 50

ggc tgg agg aga ctg gcg gag ctg gcg ggg agt cgc ggg cgc ctc cgc 368
 Gly Trp Arg Arg Leu Ala Glu Leu Ala Gly Ser Arg Gly Arg Leu Arg
 55 60 65

ctc agt tgc cta gac ctg gag cag tgt tct ctt aag gta ctg gag cct 416
 Leu Ser Cys Leu Asp Leu Glu Gln Cys Ser Leu Lys Val Leu Glu Pro
 70 75 80

gaa gga agc ccc agc ctg tgt ctg ctg aag tta atg ggt gaa aaa ggt 464
 Glu Gly Ser Pro Ser Leu Cys Leu Leu Lys Leu Met Gly Glu Lys Gly
 85 90 95 100

tgc aca gtc aca gaa ttg agt gat ttc ctg cag gct atg gaa cac act 512
 Cys Thr Val Thr Glu Leu Ser Asp Phe Leu Gln Ala Met Glu His Thr
 105 110 115

gaa gtt ctt cag ctt ctc agc ccc cca gga ata aag att act gta aac 560

Glu Val Leu Gln Leu Leu Ser Pro Pro Gly Ile Lys Ile Thr Val Asn

120 125 130

cca gag tca aag gca gtc ttg gct gga cag ttt gtg aaa ctg tgt tgc 608

Pro Glu Ser Lys Ala Val Leu Ala Gly Gln Phe Val Lys Leu Cys Cys

135 140 145

cgg gca act gga cat cct ttt gtt caa tat cag tgg ttc aaa atg aat 656

Arg Ala Thr Gly His Pro Phe Val Gln Tyr Gln Trp Phe Lys Met Asn

150 155 160

aaa gag att cca aat gga aat aca tca gag ctt att ttt aat gca gtg 704

Lys Glu Ile Pro Asn Gly Asn Thr Ser Glu Leu Ile Phe Asn Ala Val

165 170 175 180

cat gta aaa gat gca ggc ttt tat gtc tgt cga gtt aat aac aat ttc 752

His Val Lys Asp Ala Gly Phe Tyr Val Cys Arg Val Asn Asn Asn Phe

185 190 195

acc ttt gaa ttc agc cag tgg tca cag ctg gat gtt tgc gac atc cca 800

Thr Phe Glu Phe Ser Gln Trp Ser Gln Leu Asp Val Cys Asp Ile Pro

200 205 210

gag agc ttc cag aga agt gtt gat ggc gtc tct gaa tcc aag ttg caa 848

Glu Ser Phe Gln Arg Ser Val Asp Gly Val Ser Glu Ser Lys Leu Gln

215 220 225

atc tgt gtt gaa cca act tcc caa aag ctg atg cca ggc agc aca ttg 896

Ile Cys Val Glu Pro Thr Ser Gln Lys Leu Met Pro Gly Ser Thr Leu

230

235

240

gtt tta cag tgt gtt gct gga agc cct att cct cac tac cag tgg 944

Val Leu Gln Cys Val Ala Val Gly Ser Pro Ile Pro His Tyr Gln Trp

245

250

255

260

ttc aaa aat gaa tta cca tta aca cat gag acc aaa aag cta tac atg 992

Phe Lys Asn Glu Leu Pro Leu Thr His Glu Thr Lys Lys Leu Tyr Met

265

270

275

gtg cct tat gtg gat ttg gaa cac caa gga acc tac tgg tgt cat gta 1040

Val Pro Tyr Val Asp Leu Glu His Gln Gly Thr Tyr Trp Cys His Val

280

285

290

tat aat gat cga gac agt caa gat agc aag aag gta gaa atc atc ata 1088

Tyr Asn Asp Arg Asp Ser Gln Asp Ser Lys Lys Val Glu Ile Ile Ile

295

300

305

gga aga aca gat gag gca gtg gag tgc act gaa gat gaa tta aat aat 1136

Gly Arg Thr Asp Glu Ala Val Glu Cys Thr Glu Asp Glu Leu Asn Asn

310

315

320

ctt ggt cat cct gat aat aaa gag caa aca act gac cag cct ttg gcg 1184

Leu Gly His Pro Asp Asn Lys Glu Gln Thr Thr Asp Gln Pro Leu Ala

325

330

335

340

aag gac aag gtt gcc ctt ttg ata gga aat atg aat tac cgg gag cac 1232

Lys Asp Lys Val Ala Leu Leu Ile Gly Asn Met Asn Tyr Arg Glu His

345

350

355

ccc aag ctc aaa gct cct ttg gtg gat gtg tac gaa ttg act aac tta 1280

Pro Lys Leu Lys Ala Pro Leu Val Asp Val Tyr Glu Leu Thr Asn Leu

360

365

370

ctg aga cag ctg gac ttc aaa gtg gtt tca ctg ttg gat ctt act gaa 1328

Leu Arg Gln Leu Asp Phe Lys Val Val Ser Leu Leu Asp Leu Thr Glu

375

380

385

tat gag atg cgt aat gct gtg gat gag ttt tta ctc ctt tta gac aag 1376

Tyr Glu Met Arg Asn Ala Val Asp Glu Phe Leu Leu Leu Asp Lys

390

395

400

gga gta tat ggg tta tta tat tat gca gga cat ggt tat gaa aat ttt 1424

Gly Val Tyr Gly Leu Leu Tyr Tyr Ala Gly His Gly Tyr Glu Asn Phe

405

410

415

420

ggg aac agc ttc atg gtc ccc gtt gat gct cca aat cca tat agg tct 1472

Gly Asn Ser Phe Met Val Pro Val Asp Ala Pro Asn Pro Tyr Arg Ser

425

430

435

gaa aat tgt ctg tgt gta caa aat ata ctg aaa ttg atg caa gaa aaa 1520

Glu Asn Cys Leu Cys Val Gln Asn Ile Leu Lys Leu Met Gln Glu Lys

440

445

450

gaa act gga ctt aat gtg ttc tta ttg gat atg tgt agg aaa aga aat 1568

Glu Thr Gly Leu Asn Val Phe Leu Leu Asp Met Cys Arg Lys Arg Asn

455

460

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gac tac gat gat acc att cca atc ttg gat gca cta aaa gtc acc gcc 1616
 Asp Tyr Asp Asp Thr Ile Pro Ile Leu Asp Ala Leu Lys Val Thr Ala
 470 475 480

aat att gtg ttt gga tat gcc acg tgt caa gga gca gaa gct ttt gaa 1664
 Asn Ile Val Phe Gly Tyr Ala Thr Cys Gln Gly Ala Glu Ala Phe Glu
 485 490 495 500

atc cag cat tct gga ttg gca aat gga atc ttt atg aaa ttt tta aaa 1712
 Ile Gln His Ser Gly Leu Ala Asn Gly Ile Phe Met Lys Phe Leu Lys
 505 510 515

gac aga tta tta gaa gat aag aaa atc act gtg tta ctg gat gaa gtt 1760
 Asp Arg Leu Leu Glu Asp Lys Lys Ile Thr Val Leu Leu Asp Glu Val
 520 525 530

gca gaa gat atg ggt aag tgt cac ctt acc aaa ggc aaa cag gct cta 1808
 Ala Glu Asp Met Gly Lys Cys His Leu Thr Lys Gly Lys Gln Ala Leu
 535 540 545

gag att cga agt agt tta tct gag aag aga gca ctt act gat cca ata 1856
 Glu Ile Arg Ser Ser Leu Ser Glu Lys Arg Ala Leu Thr Asp Pro Ile
 550 555 560

cag gga aca gaa tat tct gct gaa tct ctt gtg cgg aat cta cag tgg 1904
 Gln Gly Thr Glu Tyr Ser Ala Glu Ser Leu Val Arg Asn Leu Gln Trp
 565 570 575 580

gcc aag gct cat gaa ctt cca gaa agt atg tgt ctt aag ttt gac tgt 1952

Ala Lys Ala His Glu Leu Pro Glu Ser Met Cys Leu Lys Phe Asp Cys

585

590

595

ggt gtt cag att caa tta gga ttt gca gct gag ttt tcc aat gtc atg 2000

Gly Val Gln Ile Gln Leu Gly Phe Ala Ala Glu Phe Ser Asn Val Met

600

605

610

atc atc tat aca agt ata gtt tac aaa cca ccg gag ata ata atg tgt 2048

Ile Ile Tyr Thr Ser Ile Val Tyr Lys Pro Pro Glu Ile Ile Met Cys

615

620

625

gat gcc tac gtt act gat ttt cca ctt gat cta gat att gat cca aaa 2096

Asp Ala Tyr Val Thr Asp Phe Pro Leu Asp Leu Asp Ile Asp Pro Lys

630

635

640

gat gca aat aaa ggc aca cct gaa gaa act ggc agc tac ttg gta tca 2144

Asp Ala Asn Lys Gly Thr Pro Glu Glu Thr Gly Ser Tyr Leu Val Ser

645

650

655

660

aag gat ctt ccc aag cat tgc ctc tat acc aga ctc agt tca ctg caa 2192

Lys Asp Leu Pro Lys His Cys Leu Tyr Thr Arg Leu Ser Ser Leu Gln

665

670

675

aaa tta aag gaa cat cta gtc ttc aca gta tgt tta tca tat cag tac 2240

Lys Leu Lys Glu His Leu Val Phe Thr Val Cys Leu Ser Tyr Gln Tyr

680

685

690

tca gga ttg gaa gat act gta gag gac aag cag gaa gtg aat gtt ggg 2288

Ser Gly Leu Glu Asp Thr Val Glu Asp Lys Gln Glu Val Asn Val Gly

695

700

705

aaa cct ctc att gct aaa tta gac atg cat cga ggt ttg gga agg aag 2336
 Lys Pro Leu Ile Ala Lys Leu Asp Met His Arg Gly Leu Gly Arg Lys

710

715

720

act tgc ttt caa act tgt ctt atg tct aat ggt cct tac cag agt tct 2384
 Thr Cys Phe Gln Thr Cys Leu Met Ser Asn Gly Pro Tyr Gln Ser Ser
 725 730 735 740

gca gcc acc tca gga gga gca ggg cat tat cac tca ttg caa gac cca 2432
 Ala Ala Thr Ser Gly Gly Ala Gly His Tyr His Ser Leu Gln Asp Pro
 745 750 755

ttc cat ggt gtt tac cat tca cat cct ggt aat cca agt aat gtt aca 2480
 Phe His Gly Val Tyr His Ser His Pro Gly Asn Pro Ser Asn Val Thr
 760 765 770

cca gca gat agc tgt cat tgc agc cgg act cca gat gca ttt att tca 2528
 Pro Ala Asp Ser Cys His Cys Ser Arg Thr Pro Asp Ala Phe Ile Ser
 775 780 785

agt ttc gct cac cat gct tca tgt cat ttt agt aga agt aat gtg cca 2576
 Ser Phe Ala His His Ala Ser Cys His Phe Ser Arg Ser Asn Val Pro
 790 795 800

gta gag aca act gat gaa ata cca ttt agt ttc tct gac agg ctc aga 2624
 Val Glu Thr Thr Asp Glu Ile Pro Phe Ser Phe Ser Asp Arg Leu Arg
 805 810 815 820

att tct gaa aaa tgacctcctt gttttgaaa gtttagataa ttttagatgc 2676
Ile Ser Glu Lys

ctgtgaaata gtactgcact tacataaagt gagacattgt gaaaaggcaa atttgtatat 2736

gtagagaaag aatagtagta actgtttcat agcaaacttc aggacttga gatgttgaaa 2796

ttacattatt taattacaga cttcctctt ct 2828

<210> 133

<211> 919

<212> PRT

<213> Homo sapiens

<400> 133

Met	Lys	Val	Ala	Arg	Phe	Gln	Lys	Ile	Pro	Asn	Gly	Glu	Asn	Glu	Thr
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Met	Ile	Pro	Val	Leu	Thr	Ser	Lys	Ala	Ser	Glu	Leu	Pro	Val	Ser
20														

Glu	Val	Ala	Ser	Ile	Leu	Gln	Ala	Asp	Leu	Gln	Asn	Gly	Leu	Asn	Lys
35															

Cys	Glu	Val	Ser	His	Arg	Arg	Ala	Phe	His	Gly	Trp	Asn	Glu	Phe	Asp
50															

Ile Ser Glu Asp Glu Pro Leu Trp Lys Lys Tyr Ile Ser Gln Phe Lys
 65 70 75 80

Asn Pro Leu Ile Met Leu Leu Ala Ser Ala Val Ile Ser Val Leu
 85 90 95

Met His Gln Phe Asp Asp Ala Val Ser Ile Thr Val Ala Ile Leu Ile
 100 105 110

Val Val Thr Val Ala Phe Val Gln Glu Tyr Arg Ser Glu Lys Ser Leu
 115 120 125

Glu Glu Leu Ser Lys Leu Val Pro Pro Glu Cys His Cys Val Arg Glu
 130 135 140

Gly Lys Leu Glu His Thr Leu Ala Arg Asp Leu Val Pro Gly Asp Thr
 145 150 155 160

Val Cys Leu Ser Val Gly Asp Arg Val Pro Ala Asp Leu Arg Leu Phe
 165 170 175

Glu Ala Val Asp Leu Ser Ile Asp Glu Ser Ser Leu Thr Gly Glu Thr
 180 185 190

Thr Pro Cys Ser Lys Val Thr Ala Pro Gln Pro Ala Ala Thr Asn Gly
 195 200 205

Asp Leu Ala Ser Arg Ser Asn Ile Ala Phe Met Gly Thr Leu Val Arg
 210 215 220

Cys Gly Lys Ala Lys Gly Val Val Ile Gly Thr Gly Glu Asn Ser Glu

225

230

235

240

Phe Gly Glu Val Phe Lys Met Met Gln Ala Glu Glu Ala Pro Lys Thr

245

250

255

Pro Leu Gln Lys Ser Met Asp Leu Leu Gly Lys Gln Leu Ser Phe Tyr

260

265

270

Ser Phe Gly Ile Ile Gly Ile Ile Met Leu Val Gly Trp Leu Leu Gly

275

280

285

Lys Asp Ile Leu Glu Met Phe Thr Ile Ser Val Ser Leu Ala Val Ala

290

295

300

Ala Ile Pro Glu Gly Leu Pro Ile Val Val Thr Val Thr Leu Ala Leu

305

310

315

320

Gly Val Met Arg Met Val Lys Lys Arg Ala Ile Val Lys Lys Leu Pro

325

330

335

Ile Val Glu Thr Leu Gly Cys Cys Asn Val Ile Cys Ser Asp Lys Thr

340

345

350

Gly Thr Leu Thr Lys Asn Glu Met Thr Val Thr His Ile Phe Thr Ser

355

360

365

Asp Gly Leu His Ala Glu Val Thr Gly Val Gly Tyr Asn Gln Phe Gly

370 375 380
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Val Ser Arg Ile Val Glu Ala Gly Cys Val Cys Asn Asp Ala Val Ile
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Arg Asn Asn Thr Leu Met Gly Lys Pro Thr Glu Gly Ala Leu Ile Ala
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Leu Ala Met Lys Met Gly Leu Asp Gly Leu Gln Gln Asp Tyr Ile Arg
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Lys Ala Glu Tyr Pro Phe Ser Ser Glu Gln Lys Trp Met Ala Val Lys
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Cys Val His Arg Thr Gln Gln Asp Arg Pro Glu Ile Cys Phe Met Lys
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Gly Ala Tyr Glu Gln Val Ile Lys Tyr Cys Thr Thr Tyr Gln Ser Lys
485 490 495

Gly Gln Thr Leu Thr Leu Thr Gln Gln Gln Arg Asp Val Tyr Gln Gln
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Glu Lys Ala Arg Met Gly Ser Ala Gly Leu Arg Val Leu Ala Leu Ala
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Ser Gly Pro Glu Leu Gly Gln Leu Thr Phe Leu Gly Leu Val Gly Ile

530 535 540

Ile Asp Pro Pro Arg Thr Gly Val Lys Glu Ala Val Thr Thr Leu Ile

545 550 555 560

Ala Ser Gly Val Ser Ile Lys Met Ile Thr Gly Asp Ser Gln Glu Thr

565 570 575

Ala Val Ala Ile Ala Ser Arg Leu Gly Leu Tyr Ser Lys Thr Ser Gln

580 585 590

Ser Val Ser Gly Glu Glu Ile Asp Ala Met Asp Val Gln Gln Leu Ser

595 600 605

Gln Ile Val Pro Lys Val Ala Val Phe Tyr Arg Ala Ser Pro Arg His

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Lys Met Lys Ile Ile Lys Ser Leu Gln Lys Asn Gly Ser Val Val Ala

625 630 635 640

Met Thr Gly Asp Gly Val Asn Asp Ala Val Ala Leu Lys Ala Ala Asp

645 650 655

Ile Gly Val Ala Met Gly Gln Thr Gly Thr Asp Val Cys Lys Glu Ala

660 665 670

Ala Asp Met Ile Leu Val Asp Asp Asp Phe Gln Thr Ile Met Ser Ala

675 680 685

Ile Glu Glu Gly Lys Gly Ile Tyr Asn Asn Ile Lys Asn Phe Val Arg

690 695 700

Phe Gln Leu Ser Thr Ser Ile Ala Ala Leu Thr Leu Ile Ser Leu Ala

705 710 715 720

Thr Leu Met Asn Phe Pro Asn Pro Leu Asn Ala Met Gln Ile Leu Trp

725 730 735

Ile Asn Ile Ile Met Asp Gly Pro Pro Ala Gln Ser Leu Gly Val Glu

740 745 750

Pro Val Asp Lys Asp Val Ile Arg Lys Pro Pro Arg Asn Trp Lys Asp

755 760 765

Ser Ile Leu Thr Lys Asn Leu Ile Leu Lys Ile Leu Val Ser Ser Ile

770 775 780

Ile Ile Val Cys Gly Thr Leu Phe Val Phe Trp Arg Glu Leu Arg Asp

785 790 795 800

Asn Val Ile Thr Pro Arg Asp Thr Thr Met Thr Phe Thr Cys Phe Val

805 810 815

Phe Phe Asp Met Phe Asn Ala Leu Ser Ser Arg Ser Gln Thr Lys Ser

820 825 830

Val Phe Glu Ile Gly Leu Cys Ser Asn Arg Met Phe Cys Tyr Ala Val

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835

840

845

Leu Gly Ser Ile Met Gly Gln Leu Leu Val Ile Tyr Phe Pro Pro Leu

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855

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Gln Lys Val Phe Gln Thr Glu Ser Leu Ser Ile Leu Asp Leu Leu Phe

865

870

875

880

Leu Leu Gly Leu Thr Ser Ser Val Cys Ile Val Ala Glu Ile Ile Lys

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gagcaggctc cgcgcctcgca ccgcgtcccc gcgcgcgcgc cctcttctcc cgaggcgcc 180

ggggcgcccc cgcgagcccc gcggctgaga cccgcagcc tggaggaggg ctgtccgggg 240

ctttggatgc tgctgcttagg ggtgggtggga gcagccgtgg gacgcgtggc cgggagcggg 300

ggtgacagcc tgggattccg ggggcttctc ttccctgtcc tcctcctctc ctctctattc 360

ccagtgtggc cgtggctgac actaaagact ttgttagccat caacccgagt gcagtttcga 420

tggaaa atg aag gtt gca cgt ttt caa aaa ata cct aat ggt gaa aat 468

Met Lys Val Ala Arg Phe Gln Lys Ile Pro Asn Gly Glu Asn

1	5	10
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gag aca atg att cct gta ttg aca tca aaa aaa gca agt gaa tta cca 516

Glu Thr Met Ile Pro Val Leu Thr Ser Lys Lys Ala Ser Glu Leu Pro

15	20	25	30
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gtc agt gaa gtt gca agc att ctc caa gct gat ctt cag aat ggt cta 564

Val Ser Glu Val Ala Ser Ile Leu Gln Ala Asp Leu Gln Asn Gly Leu

35	40	45
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Asn Lys Cys Glu Val Ser His Arg Arg Ala Phe His Gly Trp Asn Glu

50	55	60
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Phe Asp Ile Ser Glu Asp Glu Pro Leu Trp Lys Lys Tyr Ile Ser Gln			
65	70	75	
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Phe Lys Asn Pro Leu Ile Met Leu Leu Ala Ser Ala Val Ile Ser			
80	85	90	
gtt tta atg cat cag ttt gat gat gcc gtc agt atc act gtg gca ata	756		
Val Leu Met His Gln Phe Asp Asp Ala Val Ser Ile Thr Val Ala Ile			
95	100	105	110
ctt atc gtt gtt aca gtt gcc ttt gtt cag gaa tat cgt tca gaa aaa	804		
Leu Ile Val Val Thr Val Ala Phe Val Gln Glu Tyr Arg Ser Glu Lys			
115	120	125	
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Ser Leu Glu Glu Leu Ser Lys Leu Val Pro Pro Glu Cys His Cys Val			
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cgt gaa gga aaa ttg gag cat aca ctt gcc cga gac ttg gtt cca ggt	900		
Arg Glu Gly Lys Leu Glu His Thr Leu Ala Arg Asp Leu Val Pro Gly			
145	150	155	
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Asp Thr Val Cys Leu Ser Val Gly Asp Arg Val Pro Ala Asp Leu Arg			
160	165	170	
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Leu Phe Glu Ala Val Asp Leu Ser Ile Asp Glu Ser Ser Leu Thr Gly
 175 180 185 190

gag aca acg cct tgt tct aag gtg aca gct cct cag cca gct gca act 1044
 Glu Thr Thr Pro Cys Ser Lys Val Thr Ala Pro Gln Pro Ala Ala Thr
 195 200 205

aat gga gat ctt gca tcg aga agt aac att gcc ttt atg gga aca ctg 1092
 Asn Gly Asp Leu Ala Ser Arg Ser Asn Ile Ala Phe Met Gly Thr Leu
 210 215 220

gtc aga tgt ggc aaa gca aag ggt gtt gtc att gga aca gga gaa aat 1140
 Val Arg Cys Gly Lys Ala Lys Gly Val Val Ile Gly Thr Gly Glu Asn
 225 230 235

tct gaa ttt ggg gag gtt ttt aaa atg atg caa gca gaa gag gca cca 1188
 Ser Glu Phe Gly Glu Val Phe Lys Met Met Gln Ala Glu Glu Ala Pro
 240 245 250

aaa acc cct ctg cag aag agc atg gac ctc tta gga aaa caa ctt tcc 1236
 Lys Thr Pro Leu Gln Lys Ser Met Asp Leu Leu Gly Lys Gln Leu Ser
 255 260 265 270

ttt tac tcc ttt ggt ata ata gga atc atc atg ttg gtt ggc tgg tta 1284
 Phe Tyr Ser Phe Gly Ile Ile Gly Ile Ile Met Leu Val Gly Trp Leu
 275 280 285

ctg gga aaa gat atc ctg gaa atg ttt act att agt gta agt ttg gct 1332
 Leu Gly Lys Asp Ile Leu Glu Met Phe Thr Ile Ser Val Ser Leu Ala

290	295	300	
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Val Ala Ala Ile Pro Glu Gly Leu Pro Ile Val Val Thr Val Thr Leu			
305	310	315	
 gct ctt ggt gtt atg aga atg gtg aag aaa agg gcc att gtg aaa aag 1428			
Ala Leu Gly Val Met Arg Met Val Lys Lys Arg Ala Ile Val Lys Lys			
320	325	330	
 ctg cct att gtt gaa act ctg ggc tgc tgt aat gtg att tgt tca gat 1476			
Leu Pro Ile Val Glu Thr Leu Gly Cys Cys Asn Val Ile Cys Ser Asp			
335	340	345	350
 aaa act gga aca ctg acg aag aat gaa atg act gtt act cac ata ttt 1524			
Lys Thr Gly Thr Leu Thr Lys Asn Glu Met Thr Val Thr His Ile Phe			
355	360	365	
 act tca gat ggt ctg cat gct gag gtt act gga gtt ggc tat aat caa 1572			
Thr Ser Asp Gly Leu His Ala Glu Val Thr Gly Val Gly Tyr Asn Gln			
370	375	380	
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Phe Gly Glu Val Ile Val Asp Gly Asp Val Val His Gly Phe Tyr Asn			
385	390	395	
 cca gct gtt agc aga att gtt gag gcg ggc tgt gtg tgc aat gat gct 1668			
Pro Ala Val Ser Arg Ile Val Glu Ala Gly Cys Val Cys Asn Asp Ala			
400	405	410	

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Val Ile Arg Asn Asn Thr Leu Met Gly Lys Pro Thr Glu Gly Ala Leu

415 420 425 430

att gct ctt gca atg aag atg ggt ctt gat gga ctt caa caa gac tac 1764

Ile Ala Leu Ala Met Lys Met Gly Leu Asp Gly Leu Gln Gln Asp Tyr

435 440 445

atc aga aaa gct gaa tac cct ttt agc tct gag caa aag tgg atg gct 1812

Ile Arg Lys Ala Glu Tyr Pro Phe Ser Ser Glu Gln Lys Trp Met Ala

450 455 460

gtt aag tgt gta cac cga aca cag cag gac aga cca gag att tgt ttt 1860

Val Lys Cys Val His Arg Thr Gln Gln Asp Arg Pro Glu Ile Cys Phe

465 470 475

atg aaa ggt gct tac gaa caa gta att aag tac tgt act aca tac cag 1908

Met Lys Gly Ala Tyr Glu Gln Val Ile Lys Tyr Cys Thr Thr Tyr Gln

480 485 490

agc aaa ggg cag acc ttg aca ctt act cag cag cag aga gat gtg tac 1956

Ser Lys Gly Gln Thr Leu Thr Leu Thr Gln Gln Gln Arg Asp Val Tyr

495 500 505 510

caa caa gag aag gca cgc atg ggc tca gcg gga ctc aga gtt ctt gct 2004

Gln Gln Glu Lys Ala Arg Met Gly Ser Ala Gly Leu Arg Val Leu Ala

515 520 525

ttg gct tct ggt cct gaa ctg gga cag ctg aca ttt ctt ggc ttg gtg			2052
Leu Ala Ser Gly Pro Glu Leu Gly Gln Leu Thr Phe Leu Gly Leu Val			
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gga atc att gat cca cct aga act ggt gtg aaa gaa gct gtt aca aca			2100
Gly Ile Ile Asp Pro Pro Arg Thr Gly Val Lys Glu Ala Val Thr Thr			
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ctc att gcc tca gga gta tca ata aaa atg att act gga gat tca cag			2148
Leu Ile Ala Ser Gly Val Ser Ile Lys Met Ile Thr Gly Asp Ser Gln			
560	565	570	
gag act gca gtt gca atc gcc agt cgt ctg gga ttg tat tcc aaa act			2196
Glu Thr Ala Val Ala Ile Ala Ser Arg Leu Gly Leu Tyr Ser Lys Thr			
575	580	585	590
tcc cag tca gtc tca gga gaa ata gat gca atg gat gtt cag cag			2244
Ser Gln Ser Val Ser Gly Glu Glu Ile Asp Ala Met Asp Val Gln Gln			
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ctt tca caa ata gta cca aag gtt gca gta ttt tac aga gct agc cca			2292
Leu Ser Gln Ile Val Pro Lys Val Ala Val Phe Tyr Arg Ala Ser Pro			
610	615	620	
agg cac aag atg aaa att att aag tcg cta cag aag aac ggt tca gtt			2340
Arg His Lys Met Lys Ile Ile Lys Ser Leu Gln Lys Asn Gly Ser Val			
625	630	635	
gta gcc atg aca gga gat gga aat gat gca gtt gct ctg aag gct			2388

Val Ala Met Thr Gly Asp Gly Val Asn Asp Ala Val Ala Leu Lys Ala

640 645 650

gca gac att gga gtt gcg atg ggc cag act ggt aca gat gtt tgc aaa 2436

Ala Asp Ile Gly Val Ala Met Gly Gln Thr Gly Thr Asp Val Cys Lys

655 660 665 670

gag gca gca gac atg atc cta gtg gat gat gat ttt caa acc ata atg 2484

Glu Ala Ala Asp Met Ile Leu Val Asp Asp Asp Phe Gln Thr Ile Met

675 680 685

tct gca atc gaa gag ggt aaa ggg att tat aat aac att aaa aat ttc 2532

Ser Ala Ile Glu Glu Gly Lys Gly Ile Tyr Asn Asn Ile Lys Asn Phe

690 695 700

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Val Arg Phe Gln Leu Ser Thr Ser Ile Ala Ala Leu Thr Leu Ile Ser

705 710 715

ttg gct aca tta atg aac ttt cct aat cct ctc aat gcc atg cag att 2628

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720 725 730

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735 740 745 750

gta gaa cca gtg gat aaa gat gtc att cgt aaa cct cct cgc aac tgg 2724

Val Glu Pro Val Asp Lys Asp Val Ile Arg Lys Pro Pro Arg Asn Trp

755	760	765	
aaa gac agc att ttg act aaa aac ttg ata ctt aaa ata ctt gtt tca 2772			
Lys Asp Ser Ile Leu Thr Lys Asn Leu Ile Leu Lys Ile Leu Val Ser			
770	775	780	
tca ata atc att gtt tgt ggg act ttg ttt gtc ttc tgg cgt gag cta 2820			
Ser Ile Ile Ile Val Cys Gly Thr Leu Phe Val Phe Trp Arg Glu Leu			
785	790	795	
cga gac aat gtg att aca cct cga gac aca aca atg acc ttc aca tgc 2868			
Arg Asp Asn Val Ile Thr Pro Arg Asp Thr Thr Met Thr Phe Thr Cys			
800	805	810	
ttt gtg ttt ttt gac atg ttc aat gca cta agt tcc aga tcc cag acc 2916			
Phe Val Phe Phe Asp Met Phe Asn Ala Leu Ser Ser Arg Ser Gln Thr			
815	820	825	830
aag tct gtg ttt gag att gga ctc tgc agt aat aga atg ttt tgc tat 2964			
Lys Ser Val Phe Glu Ile Gly Leu Cys Ser Asn Arg Met Phe Cys Tyr			
835	840	845	
gca gtt ctt gga tcc atc atg gga caa tta cta gtt att tac ttt cct 3012			
Ala Val Leu Gly Ser Ile Met Gly Gln Leu Leu Val Ile Tyr Phe Pro			
850	855	860	
ccg ctt cag aag gtt ttt cag act gag agc cta agc ata ctg gat ctg 3060			
Pro Leu Gln Lys Val Phe Gln Thr Glu Ser Leu Ser Ile Leu Asp Leu			
865	870	875	

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 Leu Phe Leu Leu Gly Leu Thr Ser Ser Val Cys Ile Val Ala Glu Ile

880 885 890

ata aag aag gtt gaa agg agc agg gaa aag atc cag aag cat gtt agt 3156
 Ile Lys Lys Val Glu Arg Ser Arg Glu Lys Ile Gln Lys His Val Ser
 895 900 905 910

tcg aca tca tca tct ttt ctt gaa gta tgatgcataat tgcattat 3203
 Ser Thr Ser Ser Phe Leu Glu Val
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 35 40 45

Gly Leu Glu Ser Trp Ser Phe Phe Ala Val Tyr Asp Gly His Ala Gly
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Ser Gln Val Ala Lys Tyr Cys Cys Glu His Leu Leu Asp His Ile Thr
 65 70 75 80

Asn Asn Gln Asp Phe Lys Gly Ser Ala Gly Ala Pro Ser Val Glu Asn
 85 90 95

Val Lys Asn Gly Ile Arg Thr Gly Phe Leu Glu Ile Asp Glu His Met
 100 105 110

Arg Val Met Ser Glu Lys Lys His Gly Ala Asp Arg Ser Gly Ser Thr
 115 120 125

Ala Val Gly Val Leu Ile Ser Pro Gln His Thr Tyr Phe Ile Asn Cys

130 135 140

Gly Asp Ser Arg Gly Leu Leu Cys Arg Asn Arg Lys Val His Phe Phe

145 150 155 160

Thr Gln Asp His Lys Pro Ser Asn Pro Leu Glu Lys Glu Arg Ile Gln

165 170 175

Asn Ala Gly Gly Ser Val Met Ile Gln Arg Val Asn Gly Ser Leu Ala

180 185 190

Val Ser Arg Ala Leu Gly Asp Phe Asp Tyr Lys Cys Val His Gly Lys

195 200 205

Gly Pro Thr Glu Gln Leu Val Ser Pro Glu Pro Glu Val His Asp Ile

210 215 220

Glu Arg Ser Glu Glu Asp Asp Gln Phe Ile Ile Leu Ala Cys Asp Gly

225 230 235 240

Ile Trp Asp Val Met Gly Asn Glu Glu Leu Cys Asp Phe Val Arg Ser

245 250 255

Arg Leu Glu Val Thr Asp Asp Leu Glu Lys Val Cys Asn Glu Val Val

260 265 270

Asp Thr Cys Leu Tyr Lys Gly Ser Arg Asp Asn Met Ser Val Ile Leu

275 280 285

Ile Cys Phe Pro Asn Ala Pro Lys Val Ser Pro Glu Ala Val Lys Lys

290 295 300

Glu Ala Glu Leu Asp Lys Tyr Leu Glu Cys Arg Val Glu Glu Ile Ile

305 310 315 320

Lys Lys Gln Gly Glu Gly Val Pro Asp Leu Val His Val Met Arg Thr

325 330 335

Leu Ala Ser Glu Asn Ile Pro Ser Leu Pro Pro Gly Gly Glu Leu Ala

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Lys Asn Asp Asp Thr Asp Ser Thr Ser Thr Asp Asp Met Trp

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tcttagacagc tgaggcgcga aagcgatgag tcctcggttc ttccctcc ttctccggaa 180

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cgggggccccg gacgcagccc ggctccccc ctccctccg cttcccccag cctgacactgg 300

ccgcgcgtg cagcggtgac ccctcccccg gctgccgcg tcgcccgcg ggtgacccccc 360

tccccggctg ccgcgcgcgc cgccctggcc gaccaggac ctgcccgcct gcggctgctc 420

cgAACCTAGA ggATCAAGAC ATA ATG GGA GCA TTT TTA GAC AAG CCA AAG ATG 473

Met Gly Ala Phe Leu Asp Lys Pro Lys Met

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gaa aag cat aat gcc cag ggg cag ggt aat ggg ttg cga tat ggg cta 521

Glu Lys His Asn Ala Gln Gly Gln Gly Asn Gly Leu Arg Tyr Gly Leu

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agc agc atg caa ggc tgg cgt gtt gaa atg gag gat gca cat acg gct 569

Ser Ser Met Gln Gly Trp Arg Val Glu Met Glu Asp Ala His Thr Ala

30

35

40

gtg atc ggt ttg cca agt gga ctt gaa tcg tgg tca ttc ttt gct gtg 617

Val Ile Gly Leu Pro Ser Gly Leu Glu Ser Trp Ser Phe Phe Ala Val

45

50

55

tat gat ggg cat gct ggt tct cag gtt gcc aaa tac tgc tgt gag cat 665
 Tyr Asp Gly His Ala Gly Ser Gln Val Ala Lys Tyr Cys Cys Glu His

60

65

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 Leu Leu Asp His Ile Thr Asn Asn Gln Asp Phe Lys Gly Ser Ala Gly
 75 80 85 90

gca cct tct gtg gaa aat gta aag aat gga atc aga aca ggt ttt ctg 761
 Ala Pro Ser Val Glu Asn Val Lys Asn Gly Ile Arg Thr Gly Phe Leu
 95 100 105

gag att gat gaa cac atg aga gtt atg tca gag aag aaa cat ggt gca 809
 Glu Ile Asp Glu His Met Arg Val Met Ser Glu Lys Lys His Gly Ala
 110 115 120

gat aga agt ggg tca aca gct gta ggt gtc tta att tct ccc caa cat 857
 Asp Arg Ser Gly Ser Thr Ala Val Gly Val Leu Ile Ser Pro Gln His
 125 130 135

act tat ttc att aac tgt gga gac tca aga ggt tta ctt tgt agg aac 905
 Thr Tyr Phe Ile Asn Cys Gly Asp Ser Arg Gly Leu Leu Cys Arg Asn
 140 145 150

agg aaa gtt cat ttc ttc aca caa gat cac aaa cca agt aat ccg ctg 953
 Arg Lys Val His Phe Phe Thr Gln Asp His Lys Pro Ser Asn Pro Leu
 155 160 165 170

gag aaa gaa cga att cag aat gca ggt ggc tct gta atg att cag cgt 1001
 Glu Lys Glu Arg Ile Gln Asn Ala Gly Gly Ser Val Met Ile Gln Arg

175 180 185

gtg aat ggc tct ctg gct gta tcg agg gcc ctt ggg gat ttt gat tac 1049
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190 195 200

aaa tgt gtc cat gga aaa ggt cct act gag cag ctt gtc tca cca gag 1097
 Lys Cys Val His Gly Lys Gly Pro Thr Glu Gln Leu Val Ser Pro Glu

205 210 215

cct gaa gtc cat gat att gaa aga tct gaa gaa gat gat cag ttc att 1145
 Pro Glu Val His Asp Ile Glu Arg Ser Glu Glu Asp Asp Gln Phe Ile

220 225 230

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235 240 245 250

tgt gat ttt gta aga tcc aga ctt gaa gtc act gat gac ctt gag aaa 1241
 Cys Asp Phe Val Arg Ser Arg Leu Glu Val Thr Asp Asp Leu Glu Lys

255 260 265

gtt tgc aat gaa gta gtc gac acc tgt ttg tat aag gga agt cga gac 1289
 Val Cys Asn Glu Val Val Asp Thr Cys Leu Tyr Lys Gly Ser Arg Asp

270 275 280

aac atg agt gtg att ttg atc tgt ttt cca aat gca ccc aaa gta tcg 1337
 Asn Met Ser Val Ile Leu Ile Cys Phe Pro Asn Ala Pro Lys Val Ser
 285 290 295

cca gaa gca gtg aag aag gag gca gag ttg gac aag tac ctg gaa tgc 1385
 Pro Glu Ala Val Lys Lys Glu Ala Glu Leu Asp Lys Tyr Leu Glu Cys
 300 305 310

aga gta gaa gaa atc ata aag aag cag ggg gaa ggc gtc ccc gac tta 1433
 Arg Val Glu Glu Ile Ile Lys Lys Gln Gly Glu Gly Val Pro Asp Leu
 315 320 325 330

gtc cat gtg atg cgc aca tta gcg agt gag aac atc ccc agc ctc cca 1481
 Val His Val Met Arg Thr Leu Ala Ser Glu Asn Ile Pro Ser Leu Pro
 335 340 345

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 Pro Gly Gly Glu Leu Ala Ser Lys Arg Asn Val Ile Glu Ala Val Tyr
 350 355 360

aat aga ctg aat cct tac aaa aat gac gac act gac tct aca tca aca 1577
 Asn Arg Leu Asn Pro Tyr Lys Asn Asp Asp Thr Asp Ser Thr Ser Thr
 365 370 375

gat gat atg tgg taaaactgct catctagcca tggagttac cttcacctcc 1629
 Asp Asp Met Trp
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ggggatatga catgggtgag aatgattaca tcagagaact tcagcagtac aacagctagc 1749

ccagaactga tttttttt tttttgtaa atttgagact tatgtaagcg tgattcaaa 1809

ccataattcg ttttgtaaat cagactccag caatttgt tgtatgatt tttttttt 1869

taaagtgtaa ttgtccgtt acaaaatgct catatttat tatgaactgc tttaaatcac 1929

tatcaaagtt acaagaaaatg tttggcttat tgtgtgatgc aacagatata tagcccttc 1989

aagtcatgtt gtgtttggac ttggggttgg aacagggaga gcagcagcca tgcgtac 2049

acgctcaa at gtgcagatga ttatggaaaa taacctcaaa atcttacaaa gctgaacatc 2109

caaggagitta ttgaaaacta tcttaatgt tcttggtagg ggagttggca ttgttgataa 2169

agccagtc cttcatttaac tgtcttcag gatgttcctt cggtgttcc atgagtattt 2229

caggtaataa tacagtgtat tcataagaat ctcaatctt gggctaaatg cttgtttct 2289

ttgcacctct ttcaagtcc ttacatcaa ttactaattt ataagcagca gtttcctaca 2349

tatagtagga aactgccaca ttttgctat catgattggc tgggcctgct gctgttccta 2409

gtaagatatt ctgaattcca ttttatcaat aaagcttcat ttaacaaaca agaaactt 2467

<211> 358

<212> PRT

<213> Homo sapiens

<400> 137

Met Met Gln Arg Val Phe Arg Gly Lys Leu Leu Ser Asn Asp Glu Val

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Thr Ile Lys Tyr Lys Asp Glu Asp Gly Asp Leu Ile Thr Ile Phe Asp

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25

30

Ser Ser Asp Leu Ser Phe Ala Ile Gln Cys Ser Arg Ile Leu Lys Leu

35

40

45

Thr Leu Phe Val Asn Gly Gln Pro Arg Pro Leu Glu Ser Ser Gln Val

50

55

60

Lys Tyr Leu Arg Arg Glu Leu Ile Glu Leu Arg Asn Lys Val Asn Arg

65

70

75

80

Leu Leu Asp Ser Leu Glu Pro Pro Gly Glu Pro Gly Pro Ser Thr Asn

85

90

95

Ile Pro Glu Asn Asp Thr Val Asp Gly Arg Glu Glu Lys Ser Ala Ser

100

105

110

Asp Ser Ser Gly Lys Gln Ser Thr Gln Val Met Ala Ala Ser Met Ser

115

120

125

Ala Phe Asp Pro Leu Lys Asn Gln Asp Glu Ile Asn Lys Asn Val Met

130 135 140

Ser Ala Phe Gly Leu Thr Asp Asp Gln Val Ser Gly Pro Pro Ser Ala

145 150 155 160

Pro Ala Glu Asp Arg Ser Gly Thr Pro Asp Ser Ile Ala Ser Ser Ser

165 170 175

Ser Ala Ala His Pro Pro Gly Val Gln Pro Gln Gln Pro Pro Tyr Thr

180 185 190

Gly Ala Gln Thr Gln Ala Gly Gln Met Tyr Gln Gln Tyr Gln Gln Gln

195 200 205

Ala Gly Tyr Gly Ala Gln Gln Pro Gln Ala Pro Pro Gln Gln Pro Gln

210 215 220

Gln Tyr Gly Ile Gln Tyr Ser Ala Ser Tyr Ser Gln Gln Thr Gly Pro

225 230 235 240

Gln Gln Pro Gln Gln Phe Gln Gly Tyr Gly Gln Gln Pro Thr Ser Gln

245 250 255

Ala Pro Ala Pro Ala Phe Ser Gly Gln Pro Gln Gln Leu Pro Ala Gln

260 265 270

Pro Pro Gln Gln Tyr Gln Ala Ser Asn Tyr Pro Ala Gln Thr Tyr Thr

275 280 285

Ala Gln Thr Ser Gln Pro Thr Asn Tyr Thr Val Ala Pro Ala Ser Gln

290 295 300

Pro Gly Met Ala Pro Ser Gln Pro Gly Ala Tyr Gln Pro Arg Pro Gly

305 310 315 320

Phe Thr Ser Leu Pro Gly Ser Thr Met Thr Pro Pro Pro Ser Gly Pro

325 330 335

Asn Pro Tyr Ala Arg Asn Arg Pro Pro Phe Gly Gln Gly Tyr Thr Gln

340 345 350

Pro Gly Pro Gly Tyr Arg

355

<210> 138

<211> 1519

<212> DNA

<213> Homo sapiens

<220>

<221> CDS

<222> (11)..(1084)

<400> 138

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Met Met Gln Arg Val Phe Arg Gly Lys Leu Leu Ser Asn

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gat gaa gta aca ata aag tat aaa gat gaa gat gga gat ctt ata aca 97

Asp Glu Val Thr Ile Lys Tyr Lys Asp Glu Asp Gly Asp Leu Ile Thr

15

20

25

att ttt gat agt tct gac ctt tcc ttt gca att cag tgc agt agg ata 145

Ile Phe Asp Ser Ser Asp Leu Ser Phe Ala Ile Gln Cys Ser Arg Ile

30

35

40

45

ctg aaa ctg aca tta ttt gtt aat ggc cag cca aga ccc ctt gaa tca 193

Leu Lys Leu Thr Leu Phe Val Asn Gly Gln Pro Arg Pro Leu Glu Ser

50

55

60

agt cag gtg aaa tat ctc cgt cga gaa ctg ata gaa ctt cga aat aaa 241

Ser Gln Val Lys Tyr Leu Arg Arg Glu Leu Ile Glu Leu Arg Asn Lys

65

70

75

gtg aat cgt tta ttg gat agc ttg gaa cca cct gga gaa cca gga cct 289

Val Asn Arg Leu Leu Asp Ser Leu Glu Pro Pro Gly Glu Pro Gly Pro

80

85

90

tcc acc aat att cct gaa aat gat act gtg gat ggt agg gaa gaa aag 337

Ser Thr Asn Ile Pro Glu Asn Asp Thr Val Asp Gly Arg Glu Glu Lys

95

100

105

tct gct tct gat tct tct gga aaa cag tct act cag gtt atg gca gca 385

Ser Ala Ser Asp Ser Ser Gly Lys Gln Ser Thr Gln Val Met Ala Ala

110

115

120

125

agt atg tct gct ttt gat cct tta aaa aac caa gat gaa atc aat aaa		433	
Ser Met Ser Ala Phe Asp Pro Leu Lys Asn Gln Asp Glu Ile Asn Lys			
130	135	140	
aat gtt atg tca gcg ttt ggc tta aca gat gat cag gtt tca ggg cca		481	
Asn Val Met Ser Ala Phe Gly Leu Thr Asp Asp Gln Val Ser Gly Pro			
145	150	155	
ccc agt gct cct gca gaa gat cgt tca gga aca ccc gac agc att gct		529	
Pro Ser Ala Pro Ala Glu Asp Arg Ser Gly Thr Pro Asp Ser Ile Ala			
160	165	170	
tcc tcc tcc tca gca gct cac cca cca ggc gtt cag cca cag cag cca		577	
Ser Ser Ser Ala Ala His Pro Pro Gly Val Gln Pro Gln Gln Pro			
175	180	185	
cca tat aca gga gct cag act caa gca ggt cag atg tac caa cag tac		625	
Pro Tyr Thr Gly Ala Gln Thr Gln Ala Gly Gln Met Tyr Gln Gln Tyr			
190	195	200	205
cag caa cag gcc ggc tat ggt gca cag cag ccg cag gct cca cct cag		673	
Gln Gln Gln Ala Gly Tyr Gly Ala Gln Gln Pro Gln Ala Pro Pro Gln			
210	215	220	
cag cct caa cag tat ggt att cag tat tca gca agc tat agt cag cag		721	
Gln Pro Gln Gln Tyr Gly Ile Gln Tyr Ser Ala Ser Tyr Ser Gln Gln			
225	230	235	

act gga ccc caa caa cct cag cag ttc cag gga tat ggc cag caa cca 769
 Thr Gly Pro Gln Gln Pro Gln Gln Phe Gln Gly Tyr Gly Gln Gln Pro
 240 245 250

act tcc cag gca cca gct cct gcc ttt tct ggt cag cct caa caa ctg 817
 Thr Ser Gln Ala Pro Ala Pro Ala Phe Ser Gly Gln Pro Gln Gln Leu
 255 260 265

cct gct cag ccg cca cag cag tac cag gcg agc aat tat cct gca caa 865
 Pro Ala Gln Pro Pro Gln Gln Tyr Gln Ala Ser Asn Tyr Pro Ala Gln
 270 275 280 285

act tac act gcc caa act tct cag cct act aat tat act gtg gct cct 913
 Thr Tyr Thr Ala Gln Thr Ser Gln Pro Thr Asn Tyr Thr Val Ala Pro
 290 295 300

gcc tct caa cct gga atg gct cca agc caa cct ggg gcc tat caa cca 961
 Ala Ser Gln Pro Gly Met Ala Pro Ser Gln Pro Gly Ala Tyr Gln Pro
 305 310 315

aga cca ggt ttt act tca ctt cct gga agt acc atg acc cct cct cca 1009
 Arg Pro Gly Phe Thr Ser Leu Pro Gly Ser Thr Met Thr Pro Pro Pro
 320 325 330

agt ggg cct aat cct tat gcg cgt aac cgt cct ccc ttt ggt cag ggc 1057
 Ser Gly Pro Asn Pro Tyr Ala Arg Asn Arg Pro Pro Phe Gly Gln Gly
 335 340 345

tat acc caa cct gga cct ggt tat cga taaggaggt cctctacacc 1104

Tyr Thr Gln Pro Gly Pro Gly Tyr Arg

350

355

aattaatgta gctgctagct attggcctcc caaaagactc cagtactatt ttaatttcta 1164

ttgaagaagt tcagaaattt aaaaggcagag catttttat gatatcatgg ttgggtttaa 1224

ttgaaagtat aatttgctgg aacacaaaaga ccaaaatgaa agtttttcc tcctgctta 1284

aaaatgttagc agcttcttag ttactttgga acactactct tacatgtata aagtgattga 1344

cttgcatttc tagctccct tgccggagg atattaaat gctagggtga ggtttagcca 1404

tcttactttgg cttttacta ttaacatgat gtactaaagt agagcccttt gagaatacaa 1464

gatattatgt ataaaatgta acactgatga taggtaata aagatgattg aatcc 1519

<210> 139

<211> 396

<212> PRT

<213> Homo sapiens

<400> 139

Met Asn Gly Gln Leu Asp Leu Ser Gly Lys Leu Ile Val Lys Ala Gln

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10

15

Leu Gly Glu Asp Ile Arg Arg Ile Pro Ile His Asn Glu Asp Ile Thr

20

25

30

Tyr Asp Glu Leu Val Leu Met Met Gln Arg Val Phe Arg Gly Lys Leu

35 40 45

Leu Ser Asn Asp Glu Val Thr Ile Lys Tyr Lys Asp Glu Asp Gly Asp

50 55 60

Leu Ile Thr Ile Phe Asp Ser Ser Asp Leu Ser Phe Ala Ile Gln Cys

65 70 75 80

Ser Arg Ile Leu Lys Leu Thr Leu Phe Val Asn Gly Gln Pro Arg Pro

85 90 95

Leu Glu Ser Ser Gln Val Lys Tyr Leu Arg Arg Glu Leu Ile Glu Leu

100 105 110

Arg Asn Lys Val Asn Arg Leu Leu Asp Ser Leu Glu Pro Pro Gly Glu

115 120 125

Pro Gly Pro Ser Thr Asn Ile Pro Glu Asn Asp Thr Val Asp Gly Arg

130 135 140

Glu Glu Lys Ser Ala Ser Asp Ser Ser Gly Lys Gln Ser Thr Gln Val

145 150 155 160

Met Ala Ala Ser Met Ser Ala Phe Asp Pro Leu Lys Asn Gln Asp Glu

165 170 175

Ile Asn Lys Asn Val Met Ser Ala Phe Gly Leu Thr Asp Asp Gln Val

180	185	190
Ser Gly Pro Pro Ser Ala Pro Ala Glu Asp Arg Ser Gly Thr Pro Asp		
195	200	205
Ser Ile Ala Ser Ser Ser Ala Ala His Pro Pro Gly Val Gln Pro		
210	215	220
Gln Gln Pro Pro Tyr Thr Gly Ala Gln Thr Gln Ala Gly Gln Met Tyr		
225	230	235
Gln Gln Tyr Gln Gln Gln Ala Gly Tyr Gly Ala Gln Gln Pro Gln Ala		
245	250	255
Pro Pro Gln Gln Pro Gln Gln Tyr Gly Ile Gln Tyr Ser Ala Ser Tyr		
260	265	270
Ser Gln Gln Thr Gly Pro Gln Gln Pro Gln Gln Phe Gln Gly Tyr Gly		
275	280	285
Gln Gln Pro Thr Ser Gln Ala Pro Ala Pro Ala Phe Ser Gly Gln Pro		
290	295	300
Gln Gln Leu Pro Ala Gln Pro Pro Gln Gln Tyr Gln Ala Ser Asn Tyr		
305	310	315
Pro Ala Gln Thr Tyr Thr Ala Gln Thr Ser Gln Pro Thr Asn Tyr Thr		
325	330	335

Val Ala Pro Ala Ser Gln Pro Gly Met Ala Pro Ser Gln Pro Gly Ala

340 345 350

Tyr Gln Pro Arg Pro Gly Phe Thr Ser Leu Pro Gly Ser Thr Met Thr

355 360 365

Pro Pro Pro Ser Gly Pro Asn Pro Tyr Ala Arg Asn Arg Pro Pro Phe

370 375 380

Gly Gln Gly Tyr Thr Gln Pro Gly Pro Gly Tyr Arg

385 390 395

<210> 140

<211> 1641

<212> DNA

<213> Homo sapiens

<220>

<221> CDS

<222> (19)..(1206)

<400> 140

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Met Asn Gly Gln Leu Asp Leu Ser Gly Lys Leu

1 5 10

atc gtc aaa gct caa ctt ggg gag gat att cgg cga att cct att cat 99

Ile Val Lys Ala Gln Leu Gly Glu Asp Ile Arg Arg Ile Pro Ile His

15	20	25	
aat gaa gat att act tat gat gaa tta gtg cta atg atg caa cga gtt 147			
Asn Glu Asp Ile Thr Tyr Asp Glu Leu Val Leu Met Met Gln Arg Val			
30	35	40	
ttc aga gga aaa ctt ctg agt aat gat gaa gta aca ata aag tat aaa 195			
Phe Arg Gly Leu Leu Ser Asn Asp Glu Val Thr Ile Lys Tyr Lys			
45	50	55	
gat gaa gat gga gat ctt ata aca att ttt gat agt tct gac ctt tcc 243			
Asp Glu Asp Gly Asp Leu Ile Thr Ile Phe Asp Ser Ser Asp Leu Ser			
60	65	70	75
ttt gca att cag tgc agt agg ata ctg aaa ctg aca tta ttt gtt aat 291			
Phe Ala Ile Gln Cys Ser Arg Ile Leu Lys Leu Thr Leu Phe Val Asn			
80	85	90	
ggc cag cca aga ccc ctt gaa tca agt cag gtg aaa tat ctc cgt cga 339			
Gly Gln Pro Arg Pro Leu Glu Ser Ser Gln Val Lys Tyr Leu Arg Arg			
95	100	105	
gaa ctg ata gaa ctt cga aat aaa gtg aat cgt tta ttg gat agc ttg 387			
Glu Leu Ile Glu Leu Arg Asn Lys Val Asn Arg Leu Leu Asp Ser Leu			
110	115	120	
gaa cca cct gga gaa cca gga cct tcc acc aat att cct gaa aat gat 435			
Glu Pro Pro Gly Glu Pro Gly Pro Ser Thr Asn Ile Pro Glu Asn Asp			
125	130	135	

act gtg gat ggt agg gaa gaa aag tct gct tct gat tct tct gga aaa 483
 Thr Val Asp Gly Arg Glu Glu Lys Ser Ala Ser Asp Ser Ser Gly Lys
 140 145 150 155

cag tct act cag gtt atg gca gca agt atg tct gct ttt gat cct tta 531
 Gln Ser Thr Gln Val Met Ala Ala Ser Met Ser Ala Phe Asp Pro Leu
 160 165 170

aaa aac caa gat gaa atc aat aaa aat gtt atg tca gcg ttt ggc tta 579
 Lys Asn Gln Asp Glu Ile Asn Lys Asn Val Met Ser Ala Phe Gly Leu
 175 180 185

aca gat gat cag gtt tca ggg cca ccc agt gct cct gca gaa gat cgt 627
 Thr Asp Asp Gln Val Ser Gly Pro Pro Ser Ala Pro Ala Glu Asp Arg
 190 195 200

tca gga aca ccc gac agc att gct tcc tcc tca gca gct cac cca 675
 Ser Gly Thr Pro Asp Ser Ile Ala Ser Ser Ser Ala Ala His Pro
 205 210 215

cca ggc gtt cag cca cag cag cca cca tat aca gga gct cag act caa 723
 Pro Gly Val Gln Pro Gln Gln Pro Pro Tyr Thr Gly Ala Gln Thr Gln
 220 225 230 235

gca ggt cag atg tac caa cag tac cag caa cag gcc ggc tat ggt gca 771
 Ala Gly Gln Met Tyr Gln Gln Tyr Gln Gln Gln Ala Gly Tyr Gly Ala
 240 245 250

cag cag ccg cag gct cca cct cag cag cct caa cag tat ggt att cag 819
 Gln Gln Pro Gln Ala Pro Pro Gln Gln Pro Gln Gln Tyr Gly Ile Gln
 255 260 265

tat tca gca agc tat agt cag cag act gga ccc caa caa cct cag cag 867
 Tyr Ser Ala Ser Tyr Ser Gln Gln Thr Gly Pro Gln Gln Pro Gln Gln
 270 275 280

ttc cag gga tat ggc cag caa cca act tcc cag gca cca gct cct gcc 915
 Phe Gln Gly Tyr Gly Gln Gln Pro Thr Ser Gln Ala Pro Ala Pro Ala
 285 290 295

ttt tct ggt cag cct caa caa ctg cct gct cag ccg cca cag cag tac 963
 Phe Ser Gly Gln Pro Gln Gln Leu Pro Ala Gln Pro Pro Gln Gln Tyr
 300 305 310 315

cag gcg agc aat tat cct gca caa act tac act gcc caa act tct cag 1011
 Gln Ala Ser Asn Tyr Pro Ala Gln Thr Tyr Thr Ala Gln Thr Ser Gln
 320 325 330

cct act aat tat act gtg gct cct gcc tct caa cct gga atg gct cca 1059
 Pro Thr Asn Tyr Thr Val Ala Pro Ala Ser Gln Pro Gly Met Ala Pro
 335 340 345

agc caa cct ggg gcc tat caa cca aga cca ggt ttt act tca ctt cct 1107
 Ser Gln Pro Gly Ala Tyr Gln Pro Arg Pro Gly Phe Thr Ser Leu Pro
 350 355 360

gga agt acc atg acc cct cct cca agt ggg cct aat cct tat gcg cgt 1155

Gly Ser Thr Met Thr Pro Pro Pro Ser Gly Pro Asn Pro Tyr Ala Arg

365

370

375

aac cgt cct ccc ttt ggt cag ggc tat acc caa cct gga cct ggt tat 1203

Asn Arg Pro Pro Phe Gly Gln Gly Tyr Thr Gln Pro Gly Pro Gly Tyr

380

385

390

395

cga taaggaggct cctctacacc aattaatgta gctgctagct attggcctcc 1256

Arg

caaaagactc cagtactatt ttaatttgta ttgaagaagt tcagaaattt aaaagcagag 1316

catttttat gatatcattt ttgggtttaa ttgaaagtat aatttgctgg aacacaaaaga 1376

ccaaaatgaa agtttttcc tccctgctta aaaatgtgc agcttcttag ttactttgga 1436

acactactt tacatgtata aagtgattga ctgtacttcc tagttccct tgccggagg 1496

atattaaaat gctagggtga ggtttagcca tcttacttgg cttttacta ttaacatgat 1556

gtactaaagt agagccctt gagaatacaa gatattatgt ataaaatgta acactgatga 1616

tagttata aagatgattt aatcc 1641

<210> 141

<211> 323

<212> PRT

<213> Homo sapiens

<400> 141

Met Ala Phe Ser Gly Ser Gln Ala Pro Tyr Leu Ser Pro Ala Val Pro

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Phe Ser Gly Thr Ile Gln Gly Gly Leu Gln Asp Gly Leu Gln Ile Thr

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25

30

Val Asn Gly Thr Val Leu Ser Ser Ser Gly Thr Arg Phe Ala Val Asn

35

40

45

Phe Gln Thr Gly Phe Ser Gly Asn Asp Ile Ala Phe His Phe Asn Pro

50

55

60

Arg Phe Glu Asp Gly Gly Tyr Val Val Cys Asn Thr Arg Gln Asn Gly

65

70

75

80

Ser Trp Gly Pro Glu Glu Arg Lys Thr His Met Pro Phe Gln Lys Gly

85

90

95

Met Pro Phe Asp Leu Cys Phe Leu Val Gln Ser Ser Asp Phe Lys Val

100

105

110

Met Val Asn Gly Ile Leu Phe Val Gln Tyr Phe His Arg Val Pro Phe

115

120

125

His Arg Val Asp Thr Ile Ser Val Asn Gly Ser Val Gln Leu Ser Tyr

130

135

140

Ile Ser Phe Gln Pro Pro Gly Val Trp Pro Ala Asn Pro Ala Pro Ile
 145 150 155 160

Thr Gln Thr Val Ile His Thr Val Gln Ser Ala Pro Gly Gln Met Phe
 165 170 175

Ser Thr Pro Ala Ile Pro Pro Met Met Tyr Pro His Pro Ala Tyr Pro
 180 185 190

Met Pro Phe Ile Thr Thr Ile Leu Gly Gly Leu Tyr Pro Ser Lys Ser
 195 200 205

Ile Leu Leu Ser Gly Thr Val Leu Pro Ser Ala Gln Arg Phe His Ile
 210 215 220

Asn Leu Cys Ser Gly Asn His Ile Ala Phe His Leu Asn Pro Arg Phe
 225 230 235 240

Asp Glu Asn Ala Val Val Arg Asn Thr Gln Ile Asp Asn Ser Trp Gly
 245 250 255

Ser Glu Glu Arg Ser Leu Pro Arg Lys Met Pro Phe Val Arg Gly Gln
 260 265 270

Ser Phe Ser Val Trp Ile Leu Cys Glu Ala His Cys Leu Lys Val Ala
 275 280 285

Val Asp Gly Gln His Leu Phe Glu Tyr Tyr His Arg Leu Arg Asn Leu
 290 295 300

Pro Thr Ile Asn Arg Leu Glu Val Gly Gly Asp Ile Gln Leu Thr His

305

310

315

320

Val Gln Thr

<210> 142

<211> 1616

<212> DNA

<213> Homo sapiens

<220>

<221> CDS

<222> (72)..(1040)

<400> 142

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gcggcggaga g atg gcc ttc agc ggt tcc cag gct ccc tac ctg agt cca 110

Met Ala Phe Ser Gly Ser Gln Ala Pro Tyr Leu Ser Pro

1

5

10

gct gtc ccc ttt tct ggg act att caa gga ggt ctc cag gac gga ctt 158

Ala Val Pro Phe Ser Gly Thr Ile Gln Gly Gly Leu Gln Asp Gly Leu

15

20

25

cag atc act gtc aat ggg acc gtt ctc agc tcc agt gga acc agg ttt 206

Gln Ile Thr Val Asn Gly Thr Val Leu Ser Ser Ser Gly Thr Arg Phe

30

35

40

45

gct gtg aac ttt cag act ggc ttc agt gga aat gac att gcc ttc cac 254

Ala Val Asn Phe Gln Thr Gly Phe Ser Gly Asn Asp Ile Ala Phe His

50

55

60

ttc aac cct cgg ttt gaa gat gga ggg tac gtg gtg tgc aac acg agg 302

Phe Asn Pro Arg Phe Glu Asp Gly Gly Tyr Val Val Cys Asn Thr Arg

65

70

75

cag aac gga agc tgg ggg ccc gag gag agg aag aca cac atg cct ttc 350

Gln Asn Gly Ser Trp Gly Pro Glu Glu Arg Lys Thr His Met Pro Phe

80

85

90

cag aag ggg atg ccc ttt gac ctc tgc ttc ctg gtg cag agc tca gat 398

Gln Lys Gly Met Pro Phe Asp Leu Cys Phe Leu Val Gln Ser Ser Asp

95

100

105

ttc aag gtg atg gtg aac ggg atc ctc ttc gtg cag tac ttc cac cgc 446

Phe Lys Val Met Val Asn Gly Ile Leu Phe Val Gln Tyr Phe His Arg

110

115

120

125

gtg ccc ttc cac cgt gtg gac acc atc tcc gtc aat ggc tct gtg cag 494

Val Pro Phe His Arg Val Asp Thr Ile Ser Val Asn Gly Ser Val Gln

130

135

140

ctg tcc tac atc agc ttc cag cct ccc ggc gtg tgg cct gcc aac ccg 542

Leu Ser Tyr Ile Ser Phe Gln Pro Pro Gly Val Trp Pro Ala Asn Pro

145

150

155

gct ccc att acc cag aca gtc atc cac aca gtg cag agc gcc cct gga 590
 Ala Pro Ile Thr Gln Thr Val Ile His Thr Val Gln Ser Ala Pro Gly

160

165

170

cag atg ttc tct act ccc gcc atc cca cct atg atg tac ccc cac ccc 638
 Gln Met Phe Ser Thr Pro Ala Ile Pro Pro Met Met Tyr Pro His Pro
 175 180 185

gcc tat ccg atg cct ttc atc acc acc att ctg gga ggg ctg tac cca 686
 Ala Tyr Pro Met Pro Phe Ile Thr Thr Ile Leu Gly Gly Leu Tyr Pro
 190 195 200 205

tcc aag tcc atc ctc ctg tca ggc act gtc ctg ccc agt gct cag agg 734
 Ser Lys Ser Ile Leu Leu Ser Gly Thr Val Leu Pro Ser Ala Gln Arg
 210 215 220

ttc cac atc aac ctg tgc tct ggg aac cac atc gcc ttc cac ctg aac 782
 Phe His Ile Asn Leu Cys Ser Gly Asn His Ile Ala Phe His Leu Asn
 225 230 235

ccc cgt ttt gat gag aat gct gtg gtc cgc aac acc cag atc gac aac 830
 Pro Arg Phe Asp Glu Asn Ala Val Val Arg Asn Thr Gln Ile Asp Asn
 240 245 250

tcc tgg ggg tct gag gag cga agt ctg ccc cga aaa atg ccc ttc gtc 878
 Ser Trp Gly Ser Glu Glu Arg Ser Leu Pro Arg Lys Met Pro Phe Val
 255 260 265

cgt ggc cag agc ttc tca gtg tgg atc ttg tgt gaa gct cac tgc ctc 926

Arg Gly Gln Ser Phe Ser Val Trp Ile Leu Cys Glu Ala His Cys Leu

270

275

280

285

aag gtg gcc gtg gat ggt cag cac ctg ttt gaa tac tac cat cgc ctg 974

Lys Val Ala Val Asp Gly Gln His Leu Phe Glu Tyr Tyr His Arg Leu

290

295

300

agg aac ctg ccc acc atc aac aga ctg gaa gtg ggg ggc gac atc cag 1022

Arg Asn Leu Pro Thr Ile Asn Arg Leu Glu Val Gly Gly Asp Ile Gln

305

310

315

ctg acc cat gtg cag aca taggcggctt cctggccctg gggccggggg 1070

Leu Thr His Val Gln Thr

320

ctgggtgtg gggcagtctg gtcctctca tcatccccac ttcccaggcc cagctttcc 1130

aaccctgcct gggatctggg cttaatgca gaggccatgt cttgtctgg tcctgcttct 1190

ggctacagcc accctggaac ggagaaggca gctgacgggg attgccttcc tcagccgcag 1250

cagcacctgg ggctccagct gctggaatcc taccatccca ggaggcaggg acagccaggg 1310

agaggggagg agtggcagt gaagatgaag ccccatgctc agtccccctcc catccccac 1370

gcagctccac cccagtccta agccaccagc tgtctgctcc tggtgggagg tggcctcctc 1430

agccccctcct ctctgacctt taacctcaact ctcacccttgc accgtgcacc aacccttcac 1490

ccctcctgga aagcaggcct gatggcttcc cactggcctc caccacctga ccagagtgtt 1550

ctttcagag gactggctcc tttcccagtg tcctaaaaat aaagaaatga aaatgcttgt 1610

tggcac 1616

<210> 143

<211> 136

<212> PRT

<213> Homo sapiens

<400> 143

Met Ala Gly Ala Ile Ile Glu Asn Met Ser Thr Lys Lys Leu Cys Ile

1

5

10

15

Val Gly Gly Ile Leu Leu Val Phe Gln Ile Ile Ala Phe Leu Val Gly

20

25

30

Gly Leu Ile Ala Pro Gly Pro Thr Thr Ala Val Ser Tyr Met Ser Val

35

40

45

Lys Cys Val Asp Ala Arg Lys Asn His His Lys Thr Lys Trp Phe Val

50

55

60

Pro Trp Gly Pro Asn His Cys Asp Lys Ile Arg Asp Ile Glu Glu Ala

65

70

75

80

Ile Pro Arg Glu Ile Glu Ala Asn Asp Ile Val Phe Ser Val His Ile

85 90 95

Pro Leu Pro His Met Ala Leu Ser Cys Gly Phe Leu Asp Gln Arg His

100 105 110

Gly His Leu Ser Val Cys Leu Leu Thr Val Ala Phe Gly Gly Arg Phe

115 120 125

Leu Gln Pro Leu Met His Cys Val

130 135

<210> 144

<211> 1252

<212> DNA

<213> Homo sapiens

<220>

<221> CDS

<222> (225)..(632)

<400> 144

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cccagaaagg aggcgagggaa ggagggagtg tgtgagagga gggagcaaaa agtcaccct 180

aaaacattta tttcaaggag aaaagaaaaa gggggggcgc aaaa atg gct ggg gca 236

Met Ala Gly Ala

1

att ata gaa aac atg agc acc aag aag ctg tgc att gtt ggt ggg att 284

Ile Ile Glu Asn Met Ser Thr Lys Lys Leu Cys Ile Val Gly Gly Ile

5

10

15

20

ctg ctc gtg ttc caa atc atc gcc ttt ctg gtg gga ggc ttg att gct 332

Leu Leu Val Phe Gln Ile Ile Ala Phe Leu Val Gly Gly Leu Ile Ala

25

30

35

cca ggg ccc aca acg gca gtg tcc tac atg tcg gtg aaa tgt gtg gat 380

Pro Gly Pro Thr Thr Ala Val Ser Tyr Met Ser Val Lys Cys Val Asp

40

45

50

gcc cgt aag aac cat cac aag aca aaa tgg ttc gtg cct tgg gga ccc 428

Ala Arg Lys Asn His His Lys Thr Lys Trp Phe Val Pro Trp Gly Pro

55

60

65

aat cat tgt gac aag atc cga gac att gaa gag gca att cca agg gaa 476

Asn His Cys Asp Lys Ile Arg Asp Ile Glu Glu Ala Ile Pro Arg Glu

70

75

80

att gaa gcc aat gac atc gtg ttt tct gtt cac att ccc ctc ccc cac 524

Ile Glu Ala Asn Asp Ile Val Phe Ser Val His Ile Pro Leu Pro His

85

90

95

100

atg gct ctt agc tgt ggt ttc ttg gac cag cgg cat gga cat ttg tca 572
 Met Ala Leu Ser Cys Gly Phe Leu Asp Gln Arg His Gly His Leu Ser
 105 110 115

gtt tgc ctt ctg acg gta gct ttt gga gga aga ttc ctg cag cca cta 620
 Val Cys Leu Leu Thr Val Ala Phe Gly Gly Arg Phe Leu Gln Pro Leu
 120 125 130

atg cat tgt gta tgataacaaa aactctggta tgacacattt tctgtgatca 672
 Met His Cys Val
 135

ttgttaatta gtgacatagt aacatctgta gcagctggtt agtaaacctc atgtgggggt 732

ggggtgtgggg tgcatttcattt gggggatggt ttggccgaa tggggagtgg aatatttgac 792

attttcctg ttttaaattc taggatagat tttaacatcc tttcggtcc cagtccaagg 852

taggctggtg tcatagtctt ctcactccta atccatgacc actgttttt tcctatttat 912

atcaccaggt agcctactga gttaatattt aagttgtcaa tagataagtg tccctgttt 972

gtggcataat ataactgaat ttcatgagaa gatttattcc accagggta tttcagcttt 1032

gaaaccaaattt ctgtgtatct aatactaacc aatctgttgg atgtgggttt taaaaaatgt 1092

ttgctaaact acccaagtaa gatttactgt attaaatggc cttcggtct gaaaagcttt 1152

tttaacctct tgcttaaat gcgtttattt ttgataagat acttcaaata gcctccaaaa 1212

gttagatcc aatcactaa ataaacctgt atgtatatgc

1252

<210> 145

<211> 468

<212> PRT

<213> Homo sapiens

<400> 145

Met Pro Val Arg Thr Ile Thr Arg Gln Asn Gly Ser Cys Leu Gly Asp

1

5

10

15

Pro Ile Ile Val Thr Arg Ser Glu Thr Leu Lys Arg Gln Phe Gln Phe

20

25

30

Met Leu Phe Ile Leu Gln Leu Asp Ile Ala Phe Lys Leu Asn Asn Gln

35

40

45

Ile Arg Glu Asn Ala Glu Val Ser Met Asp Val Ser Leu Ala Tyr Arg

50

55

60

Asp Asp Ala Phe Ala Glu Trp Thr Glu Met Ala His Glu Arg Val Pro

65

70

75

80

Arg Lys Leu Lys Cys Thr Phe Thr Ser Pro Lys Thr Pro Glu His Glu

85

90

95

Gly Arg Tyr Tyr Glu Cys Asp Val Leu Pro Phe Met Glu Ile Gly Ser

100	105	110
Val Ala His Lys Phe Tyr Leu Leu Asn Ile Arg Leu Pro Val Asn Glu		
115	120	125
Lys Lys Lys Ile Asn Val Gly Ile Gly Glu Ile Lys Asp Ile Arg Leu		
130	135	140
Val Gly Ile His Gln Asn Gly Gly Phe Thr Lys Val Trp Phe Ala Met		
145	150	155
160		
Lys Thr Phe Leu Thr Pro Ser Ile Phe Ile Ile Met Val Trp Tyr Trp		
165	170	175
Arg Arg Ile Thr Met Met Ser Arg Pro Pro Val Leu Leu Glu Lys Val		
180	185	190
Ile Phe Ala Leu Gly Ile Ser Met Thr Phe Ile Asn Ile Pro Val Glu		
195	200	205
Trp Phe Ser Ile Gly Phe Asp Trp Thr Trp Met Leu Leu Phe Gly Asp		
210	215	220
Ile Arg Gln Gly Ile Phe Tyr Ala Met Leu Leu Ser Phe Trp Ile Ile		
225	230	235
240		
Phe Cys Gly Glu His Met Met Asp Gln His Glu Arg Asn His Ile Ala		
245	250	255

Gly Tyr Trp Lys Gln Val Gly Pro Ile Ala Val Gly Ser Phe Cys Leu

260	265	270
-----	-----	-----

Phe Ile Phe Asp Met Cys Glu Arg Gly Val Gln Leu Thr Asn Pro Phe

275	280	285
-----	-----	-----

Tyr Ser Ile Trp Thr Asp Ile Gly Thr Glu Leu Ala Met Ala Phe

290	295	300
-----	-----	-----

Ile Ile Val Ala Gly Ile Cys Leu Cys Leu Tyr Phe Leu Phe Leu Cys

305	310	315	320
-----	-----	-----	-----

Phe Met Val Phe Gln Val Phe Arg Asn Ile Ser Gly Lys Gln Ser Ser

325	330	335
-----	-----	-----

Leu Pro Ala Met Ser Lys Val Arg Arg Leu His Tyr Glu Gly Leu Ile

340	345	350
-----	-----	-----

Phe Arg Phe Lys Phe Leu Met Leu Ile Thr Leu Ala Cys Ala Ala Met

355	360	365
-----	-----	-----

Thr Val Ile Phe Phe Ile Val Ser Gln Val Thr Glu Gly His Trp Lys

370	375	380
-----	-----	-----

Trp Gly Gly Val Thr Val Gln Val Asn Ser Ala Phe Phe Thr Gly Ile

385	390	395	400
-----	-----	-----	-----

Tyr Gly Met Trp Asn Leu Tyr Val Phe Ala Leu Met Phe Leu Tyr Ala

405	410	415
-----	-----	-----

Pro Ser His Lys Asn Tyr Gly Glu Asp Gln Ser Asn Gly Met Gln Leu

420

425

430

Pro Cys Lys Ser Arg Glu Asp Cys Ala Leu Phe Val Ser Glu Leu Tyr

435

440

445

Gln Glu Leu Phe Ser Ala Ser Lys Tyr Ser Phe Ile Asn Asp Asn Ala

450

455

460

Ala Ser Gly Ile

465

<210> 146

<211> 1943

<212> DNA

<213> Homo sapiens

<220>

<221> CDS

<222> (379)..(1782)

<400> 146

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ggcgccttcc gtcccggtcc catcctcgcc gcgctccagc acctctgaag ttttgagcg 120

cccagaaagg aggcgaggaa ggagggagtg tgtgagagga gggagcaaaa agctcaccct 180

aaaacattta tttcaaggag aaaagaaaaa gggggggcgc aaaaatggct gggcaatta 240

tagaaaacat gagcaccaag aagctgtgca ttgttggtgg gattctgctc gtgttccaaa 300

tcatgcctt tctggtgga ggcttgattt ctccagggcc cacaacggca gtgcctaca 360

tgtcggtgaa atgtgtgg atg ccc gta aga acc atc aca aga caa aat ggt 411

Met Pro Val Arg Thr Ile Thr Arg Gln Asn Gly

1	5	10
---	---	----

tcg tgc ctt ggg gac cca atc att gtg aca aga tcc gag aca ttg aag 459

Ser Cys Leu Gly Asp Pro Ile Ile Val Thr Arg Ser Glu Thr Leu Lys

15	20	25
----	----	----

agg caa ttc caa ttc atg ctg ttt atc ctg cag ctg gac att gcc ttc 507

Arg Gln Phe Gln Phe Met Leu Phe Ile Leu Gln Leu Asp Ile Ala Phe

30	35	40
----	----	----

aag cta aac aac caa atc aga gaa aat gca gaa gtc tcc atg gac gtt 555

Lys Leu Asn Asn Gln Ile Arg Glu Asn Ala Glu Val Ser Met Asp Val

45	50	55
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tcc ctg gct tac cgt gat gac gcg ttt gct gag tgg act gaa atg gcc 603

Ser Leu Ala Tyr Arg Asp Asp Ala Phe Ala Glu Trp Thr Glu Met Ala

60	65	70	75
----	----	----	----

cat gaa aga gta cca cgg aaa ctc aaa tgc acc ttc aca tct ccc aag 651

His Glu Arg Val Pro Arg Lys Leu Lys Cys Thr Phe Thr Ser Pro Lys

80	85	90	
act cca gag cat gag ggc cgt tac tat gaa tgt gat gtc ctt cct ttc 699			
Thr Pro Glu His Glu Gly Arg Tyr Tyr Glu Cys Asp Val Leu Pro Phe			
95	100	105	
atg gaa att ggg tct gtg gcc cat aag ttt tac ctt tta aac atc cgg 747			
Met Glu Ile Gly Ser Val Ala His Lys Phe Tyr Leu Leu Asn Ile Arg			
110	115	120	
ctg cct gtg aat gag aag aag aaa atc aat gtg gga att ggg gag ata 795			
Leu Pro Val Asn Glu Lys Lys Ile Asn Val Gly Ile Gly Glu Ile			
125	130	135	
aag gat atc cgg ttg gtg ggg atc cac caa aat gga ggc ttc acc aag 843			
Lys Asp Ile Arg Leu Val Gly Ile His Gln Asn Gly Gly Phe Thr Lys			
140	145	150	155
gtg tgg ttt gcc atg aag acc ttc ctt acg ccc agc atc ttc atc att 891			
Val Trp Phe Ala Met Lys Thr Phe Leu Thr Pro Ser Ile Phe Ile Ile			
160	165	170	
atg gtg tgg tat tgg agg agg atc acc atg atg tcc cga ccc cca gtg 939			
Met Val Trp Tyr Trp Arg Arg Ile Thr Met Met Ser Arg Pro Pro Val			
175	180	185	
ctt ctg gaa aaa gtc atc ttt gcc ctt ggg att tcc atg acc ttt atc 987			
Leu Leu Glu Lys Val Ile Phe Ala Leu Gly Ile Ser Met Thr Phe Ile			
190	195	200	

aat atc cca gtg gaa tgg ttt tcc atc ggg ttt gac tgg acc tgg atg 1035

Asn Ile Pro Val Glu Trp Phe Ser Ile Gly Phe Asp Trp Thr Trp Met

205

210

215

ctg ctg ttt ggt gac atc cga cag ggc atc ttc tat gcg atg ctt ctg 1083

Leu Leu Phe Gly Asp Ile Arg Gln Gly Ile Phe Tyr Ala Met Leu Leu

220

225

230

235

tcc ttc tgg atc atc ttc tgt ggc gag cac atg atg gat cag cac gag 1131

Ser Phe Trp Ile Ile Phe Cys Gly Glu His Met Met Asp Gln His Glu

240

245

250

cgg aac cac atc gca ggg tat tgg aag caa gtc gga ccc att gcc gtt 1179

Arg Asn His Ile Ala Gly Tyr Trp Lys Gln Val Gly Pro Ile Ala Val

255

260

265

ggc tcc ttc tgc ctc ttc ata ttt gac atg tgt gag aga ggg gta caa 1227

Gly Ser Phe Cys Leu Phe Ile Phe Asp Met Cys Glu Arg Gly Val Gln

270

275

280

ctc acg aat ccc ttc tac agt atc tgg act aca gac att gga aca gag 1275

Leu Thr Asn Pro Phe Tyr Ser Ile Trp Thr Thr Asp Ile Gly Thr Glu

285

290

295

ctg gcc atg gcc ttc atc atc gtg gct gga atc tgc ctc tgc ctc tac 1323

Leu Ala Met Ala Phe Ile Ile Val Ala Gly Ile Cys Leu Cys Leu Tyr

300

305

310

315

ttc ctg ttt cta tgc ttc atg gta ttt cag gtg ttt cgg aac atc agt			1371
Phe Leu Phe Leu Cys Phe Met Val Phe Gln Val Phe Arg Asn Ile Ser			
320	325	330	
ggg aag cag tcc agc ctg cca gct atg agc aaa gtc cgg cgg cta cac			1419
Gly Lys Gln Ser Ser Leu Pro Ala Met Ser Lys Val Arg Arg Leu His			
335	340	345	
tat gag ggg cta att ttt agg ttc aag ttc ctc atg ctt atc acc ttg			1467
Tyr Glu Gly Leu Ile Phe Arg Phe Lys Phe Leu Met Leu Ile Thr Leu			
350	355	360	
gcc tgc gct gcc atg act gtc atc ttc ttc atc gtt agt cag gta acg			1515
Ala Cys Ala Ala Met Thr Val Ile Phe Phe Ile Val Ser Gln Val Thr			
365	370	375	
gaa ggc cat tgg aaa tgg ggc ggc gtc aca gtc caa gtg aac agt gcc			1563
Glu Gly His Trp Lys Trp Gly Gly Val Thr Val Gln Val Asn Ser Ala			
380	385	390	395
ttt ttc aca ggc atc tat ggg atg tgg aat ctg tat gtc ttt gct ctg			1611
Phe Phe Thr Gly Ile Tyr Gly Met Trp Asn Leu Tyr Val Phe Ala Leu			
400	405	410	
atg ttc ttg tat gca cca tcc cat aaa aac tat gga gaa gac cag tcc			1659
Met Phe Leu Tyr Ala Pro Ser His Lys Asn Tyr Gly Glu Asp Gln Ser			
415	420	425	
aat gga atg caa ctc cca tgt aaa tcg agg gaa gat tgt gct ttg ttt			1707

特2001-254018

Asn Gly Met Gln Leu Pro Cys Lys Ser Arg Glu Asp Cys Ala Leu Phe

430

435

440

gtt tcg gaa ctt tat caa gaa ttg ttc agc gct tcg aaa tat tcc ttc 1755

Val Ser Glu Leu Tyr Gln Glu Leu Phe Ser Ala Ser Lys Tyr Ser Phe

445

450

455

atc aat gac aac gca gct tct ggt att tgagtcaaca aggcaacaca 1802

Ile Asn Asp Asn Ala Ala Ser Gly Ile

460

465

tgttatcag ctttgcat ttgcagttgtca cagtcacatt gattgtactt gtatacgcac 1862

acaatacac tcatttagcc ttatctcaa aatgttaat ataaggaaaa aagcgtcaac 1922

aataaatatt ctttgagttat 1943

<210> 147

<211> 460

<212> PRT

<213> Homo sapiens

<400> 147

Met Pro Val Arg Thr Ile Thr Arg Gln Asn Gly Ser Cys Leu Gly Asp

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Pro Ile Ile Val Thr Arg Ser Glu Thr Leu Lys Arg Gln Phe Gln Gly

20

25

30

Lys Leu Lys Pro Met Thr Ser Cys Phe Leu Phe Thr Phe Pro Ser Pro

35 40 45

Met Asp Val Ser Leu Ala Tyr Arg Asp Asp Ala Phe Ala Glu Trp Thr

50 55 60

Glu Met Ala His Glu Arg Val Pro Arg Lys Leu Lys Cys Thr Phe Thr

65 70 75 80

Ser Pro Lys Thr Pro Glu His Glu Gly Arg Tyr Tyr Glu Cys Asp Val

85 90 95

Leu Pro Phe Met Glu Ile Gly Ser Val Ala His Lys Phe Tyr Leu Leu

100 105 110

Asn Ile Arg Leu Pro Val Asn Glu Lys Lys Ile Asn Val Gly Ile

115 120 125

Gly Glu Ile Lys Asp Ile Arg Leu Val Gly Ile His Gln Asn Gly Gly

130 135 140

Phe Thr Lys Val Trp Phe Ala Met Lys Thr Phe Leu Thr Pro Ser Ile

145 150 155 160

Phe Ile Ile Met Val Trp Tyr Trp Arg Arg Ile Thr Met Met Ser Arg

165 170 175

Pro Pro Val Leu Leu Glu Lys Val Ile Phe Ala Leu Gly Ile Ser Met

180	185	190
Thr Phe Ile Asn Ile Pro Val Glu Trp Phe Ser Ile Gly Phe Asp Trp		
195	200	205
Thr Trp Met Leu Leu Phe Gly Asp Ile Arg Gln Gly Ile Phe Tyr Ala		
210	215	220
Met Leu Leu Ser Phe Trp Ile Ile Phe Cys Gly Glu His Met Met Asp		
225	230	235
Gln His Glu Arg Asn His Ile Ala Gly Tyr Trp Lys Gln Val Gly Pro		
245	250	255
Ile Ala Val Gly Ser Phe Cys Leu Phe Ile Phe Asp Met Cys Glu Arg		
260	265	270
Gly Val Gln Leu Thr Asn Pro Phe Tyr Ser Ile Trp Thr Thr Asp Ile		
275	280	285
Gly Thr Glu Leu Ala Met Ala Phe Ile Ile Val Ala Gly Ile Cys Leu		
290	295	300
Cys Leu Tyr Phe Leu Phe Leu Cys Phe Met Val Phe Gln Val Phe Arg		
305	310	315
Asn Ile Ser Gly Lys Gln Ser Ser Leu Pro Ala Met Ser Lys Val Arg		
325	330	335

Arg Leu His Tyr Glu Gly Leu Ile Phe Arg Phe Lys Phe Leu Met Leu

340 345 350

Ile Thr Leu Ala Cys Ala Ala Met Thr Val Ile Phe Phe Ile Val Ser

355 360 365

Gln Val Thr Glu Gly His Trp Lys Trp Gly Gly Ile Thr Val Gln Val

370 375 380

Asn Ser Ala Phe Phe Thr Gly Ile Tyr Gly Met Trp Asn Leu Tyr Val

385 390 395 400

Phe Ala Leu Met Phe Leu Tyr Ala Pro Ser His Lys Asn Tyr Gly Glu

405 410 415

Asp Gln Ser Asn Gly Met Gln Leu Pro Cys Lys Ser Arg Glu Asp Cys

420 425 430

Ala Leu Phe Val Ser Glu Leu Tyr Gln Glu Leu Phe Ser Ala Ser Lys

435 440 445

Tyr Ser Phe Ile Asn Asp Asn Ala Ala Ser Gly Ile

450 455 460

<210> 148

<211> 1919

<212> DNA

<213> Homo sapiens

<220>

<221> CDS

<222> (376)..(1755)

<400> 148

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gccttccgtc ccggccccat cctcgccgCG ctccAGCACC tctGAAGTTT tgcaGCGCCC 120

agaaaaggagg cgaggaagga gggagtgtat gagaggagg AGCAAAAAGC tcaccctaaa 180

acattttttt caaggagaaaa agaaaaagggg gggcgcaaa aatggctggg gcaattata 240

aaaacatgag caccaagaag ctgtgcattt ttggggat tctgctcgtt ttccaaatca 300

tcgcctttct ggtgggaggc ttgattgctc cagggcccac aacggcagtG tcctacatgt 360

cggtgaaatg tgtgg atg ccc gta aga acc atc aca aga caa aat ggt tcg 411

Met Pro Val Arg Thr Ile Thr Arg Gln Asn Gly Ser

1

5

10

tgc ctt ggg gac cca atc att gtg aca aga tcc gag aca ttg aag agg 459

Cys Leu Gly Asp Pro Ile Ile Val Thr Arg Ser Glu Thr Leu Lys Arg

15

20

25

caa ttc caa ggg aaa ttg aag cca atg aca tcg tgt ttt ctg ttc aca 507

Gln Phe Gln Gly Lys Leu Lys Pro Met Thr Ser Cys Phe Leu Phe Thr

30

35

40

ttc ccc tcc ccc atg gac gtt tcc ctg gct tac cgt gat gac gcg ttt 555
 Phe Pro Ser Pro Met Asp Val Ser Leu Ala Tyr Arg Asp Asp Ala Phe
 45 50 55 60

gct gag tgg act gaa atg gcc cat gaa aga gta cca cgg aaa ctc aaa 603
 Ala Glu Trp Thr Glu Met Ala His Glu Arg Val Pro Arg Lys Leu Lys
 65 70 75

tgc acc ttc aca tct ccc aag act cca gag cat gag ggc cgt tac tat 651
 Cys Thr Phe Thr Ser Pro Lys Thr Pro Glu His Glu Gly Arg Tyr Tyr
 80 85 90

gaa tgt gat gtc ctt cct ttc atg gaa att ggg tct gtg gcc cat aag 699
 Glu Cys Asp Val Leu Pro Phe Met Glu Ile Gly Ser Val Ala His Lys
 95 100 105

ttt tac ctt tta aac atc cgg ctg cct gtg aat gag aag aag aaa atc 747
 Phe Tyr Leu Leu Asn Ile Arg Leu Pro Val Asn Glu Lys Lys Ile
 110 115 120

aat gtg gga att ggg gag ata aag gat atc cgg ttg gtg ggg atc cac 795
 Asn Val Gly Ile Gly Glu Ile Lys Asp Ile Arg Leu Val Gly Ile His
 125 130 135 140

caa aat gga ggc ttc acc aag gtg tgg ttt gcc atg aag acc ttc ctt 843
 Gln Asn Gly Gly Phe Thr Lys Val Trp Phe Ala Met Lys Thr Phe Leu
 145 150 155

acg ccc agc atc ttc atc att atg gtg tgg tat tgg agg agg atc acc 891
 Thr Pro Ser Ile Phe Ile Ile Met Val Trp Tyr Trp Arg Arg Ile Thr
 160 165 170

atg atg tcc cga ccc cca gtg ctt ctg gaa aaa gtc atc ttt gcc ctt 939
 Met Met Ser Arg Pro Pro Val Leu Leu Glu Lys Val Ile Phe Ala Leu
 175 180 185

ggg att tcc atg acc ttt atc aat atc cca gtg gaa tgg ttt tcc atc 987
 Gly Ile Ser Met Thr Phe Ile Asn Ile Pro Val Glu Trp Phe Ser Ile
 190 195 200

ggg ttt gac tgg acc tgg atg ctg ctg ttt ggt gac atc cga cag ggc 1035
 Gly Phe Asp Trp Thr Trp Met Leu Leu Phe Gly Asp Ile Arg Gln Gly
 205 210 215 220

atc ttc tat gcg atg ctt ctg tcc ttc tgg atc atc ttc tgt ggc gag 1083
 Ile Phe Tyr Ala Met Leu Leu Ser Phe Trp Ile Ile Phe Cys Gly Glu
 225 230 235

cac atg atg gat cag cac gag cgg aac cac atc gca ggg tat tgg aag 1131
 His Met Met Asp Gln His Glu Arg Asn His Ile Ala Gly Tyr Trp Lys
 240 245 250

caa gtc gga ccc att gcc gtt ggc tcc ttc tgc ctc ttc ata ttt gac 1179
 Gln Val Gly Pro Ile Ala Val Gly Ser Phe Cys Leu Phe Ile Phe Asp
 255 260 265

atg tgt gag aga ggg gta caa ctc acg aat ccc ttc tac agt atc tgg 1227

Met Cys Glu Arg Gly Val Gln Leu Thr Asn Pro Phe Tyr Ser Ile Trp

270

275

280

act aca gac att gga aca gag ctg gcc atg gcc ttc atc atc gtg gct 1275

Thr Thr Asp Ile Gly Thr Glu Leu Ala Met Ala Phe Ile Ile Val Ala

285

290

295

300

gga atc tgc ctc tgc ctc tac ttc ctg ttt cta tgc ttc atg gta ttt 1323

Gly Ile Cys Leu Cys Leu Tyr Phe Leu Phe Leu Cys Phe Met Val Phe

305

310

315

cag gtg ttt cgg aac atc agt ggg aag cag tcc agc ctg cca gct atg 1371

Gln Val Phe Arg Asn Ile Ser Gly Lys Gln Ser Ser Leu Pro Ala Met

320

325

330

agc aaa gtc cgg cgg cta cac tat gag ggg cta att ttt agg ttc aag 1419

Ser Lys Val Arg Arg Leu His Tyr Glu Gly Leu Ile Phe Arg Phe Lys

335

340

345

ttc ctc atg ctt atc acc ttg gcc tgc gct gcc atg act gtc atc ttc 1467

Phe Leu Met Leu Ile Thr Leu Ala Cys Ala Ala Met Thr Val Ile Phe

350

355

360

ttc atc gtt agt cag gta acg gaa ggc cat tgg aaa tgg ggc ggc atc 1515

Phe Ile Val Ser Gln Val Thr Glu Gly His Trp Lys Trp Gly Gly Ile

365

370

375

380

aca gtc caa gtg aac agt gcc ttt ttc aca ggc atc tat ggg atg tgg 1563

Thr Val Gln Val Asn Ser Ala Phe Phe Thr Gly Ile Tyr Gly Met Trp

385

390

395

aat ctg tat gtc ttt gct ctg atg ttc ttg tat gca cca tcc cat aaa 1611

Asn Leu Tyr Val Phe Ala Leu Met Phe Leu Tyr Ala Pro Ser His Lys

400

405

410

aac tat gga gaa gac cag tcc aat gga atg caa ctc cca tgt aaa tcg 1659

Asn Tyr Gly Glu Asp Gln Ser Asn Gly Met Gln Leu Pro Cys Lys Ser

415

420

425

agg gaa gat tgt gct ttg ttt gtt tcg gaa ctt tat caa gaa ttg ttc 1707

Arg Glu Asp Cys Ala Leu Phe Val Ser Glu Leu Tyr Gln Glu Leu Phe

430

435

440

agc gct tcg aaa tat tcc ttc atc aat gac aac gca gct tct ggt att 1755

Ser Ala Ser Lys Tyr Ser Phe Ile Asn Asp Asn Ala Ala Ser Gly Ile

445

450

455

460

tgagtcaaca aggcaacaca tgtttatcag ctttgcattt gcagttgtca cagtcacatt 1815

gattgtactt gtatacgcac acaaatacac tcatttagcc tttatctcaa aatgttaaat 1875

ataaggaaaa aagcgtcaac aataaatatt cttttagtat tgtc 1919

<210> 149

<211> 183

<212> PRT

<213> Homo sapiens

<400> 149

Met Lys Leu Leu Ser Leu Val Ala Val Val Gly Cys Leu Leu Val Pro

1

5

10

15

Pro Ala Glu Ala Asn Lys Ser Ser Glu Asp Ile Arg Cys Lys Cys Ile

20

25

30

Cys Pro Pro Tyr Arg Asn Ile Ser Gly His Ile Tyr Asn Gln Asn Val

35

40

45

Ser Gln Lys Asp Cys Asn Cys Leu His Val Val Glu Pro Met Pro Val

50

55

60

Pro Gly His Asp Val Glu Ala Tyr Cys Leu Leu Cys Glu Cys Arg Tyr

65

70

75

80

Glu Glu Arg Ser Thr Thr Ile Lys Val Ile Ile Val Ile Tyr Leu

85

90

95

Ser Val Val Gly Ala Leu Leu Leu Tyr Met Ala Phe Leu Met Leu Val

100

105

110

Asp Pro Leu Ile Arg Lys Pro Asp Ala Tyr Thr Glu Gln Leu His Asn

115

120

125

Glu Glu Glu Asn Glu Asp Ala Arg Ser Met Ala Ala Ala Ala Ser

130

135

140

Leu Gly Gly Pro Arg Ala Asn Thr Val Leu Glu Arg Val Glu Gly Ala
 145 150 155 160

Gln Gln Arg Trp Lys Leu Gln Val Gln Glu Gln Arg Lys Thr Val Phe
 165 170 175

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atg aag ctc tta tct ttg gtg gct gtg gtc ggg tgt ttg ctg gtg ccc 167
 Met Lys Leu Leu Ser Leu Val Ala Val Val Gly Cys Leu Leu Val Pro
 1 5 10 15

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Pro Ala Glu Ala Asn Lys Ser Ser Glu Asp Ile Arg Cys Lys Cys Ile

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tgt cca cct tat aga aac atc agt ggg cac att tac aac cag aat gta 263

Cys Pro Pro Tyr Arg Asn Ile Ser Gly His Ile Tyr Asn Gln Asn Val

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tcc cag aag gac tgc aac tgc ctg cac gtg gtg gag ccc atg cca gtg 311

Ser Gln Lys Asp Cys Asn Cys Leu His Val Val Glu Pro Met Pro Val

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cct ggc cat gac gtg gag gcc tac tgc ctg ctg tgc gag tgc agg tac 359

Pro Gly His Asp Val Glu Ala Tyr Cys Leu Leu Cys Glu Cys Arg Tyr

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Glu Glu Arg Ser Thr Thr Ile Lys Val Ile Ile Val Ile Tyr Leu

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tcc gtg gtg ggt gcc ctg ttg ctc tac atg gcc ttc ctg atg ctg gtg 455

Ser Val Val Gly Ala Leu Leu Leu Tyr Met Ala Phe Leu Met Leu Val

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Asp Pro Leu Ile Arg Lys Pro Asp Ala Tyr Thr Glu Gln Leu His Asn

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Glu Glu Glu Asn Glu Asp Ala Arg Ser Met Ala Ala Ala Ala Ser

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Leu Gly Gly Pro Arg Ala Asn Thr Val Leu Glu Arg Val Glu Gly Ala

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cag cag cgg tgg aag ctg cag gtg cag gag cag cggtt aag aca gtc ttc 647

Gln Gln Arg Trp Lys Leu Gln Val Gln Glu Gln Arg Lys Thr Val Phe

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Asp Arg His Lys Met Leu Ser

180

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atctgaacac cacagccccct gtacttgggt tgcctttgt ccctgaacctt cgttgtacca 1478

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aggctgtcgg ttcggaacat gtctccaccc accccaccct ctgtggctcc aggcttcatt 180

ctcccccatc c atg gat aac cca ggg cct tcg ctc cgt ggt gcc ttt ggc 230
Met Asp Asn Pro Gly Pro Ser Leu Arg Gly Ala Phe Gly

1 5 10

att cta ggt gcc ttg gaa agg gac agg ctg acc cac ctg aaa cac aag 278
Ile Leu Gly Ala Leu Glu Arg Asp Arg Leu Thr His Leu Lys His Lys

15 20 25

ctg ggg agt ctg tgt tca ggc agc cag gag tca aag ctt ctc cat gcc 326
Leu Gly Ser Leu Cys Ser Gly Ser Gln Glu Ser Lys Leu Leu His Ala
30 35 40 45

atg gta ctc ctg gct ctg ggc cag gac acg gag gcc agg gtc tct ctg 374
Met Val Leu Leu Ala Leu Gly Gln Asp Thr Glu Ala Arg Val Ser Leu
50 55 60

gag tcc ttg aag atg aac aca gta gcc cag ctg gta gcc cac cag tgg 422
Glu Ser Leu Lys Met Asn Thr Val Ala Gln Leu Val Ala His Gln Trp
65 70 75

gca gac atg gag acc aca gag ggc cct gag gag cct cca gac ttg tcc 470
Ala Asp Met Glu Thr Thr Glu Gly Pro Glu Glu Pro Pro Asp Leu Ser
80 85 90

tgg acg gtg gct cgc ctg tac cac ctg ctg gct gag gag aac ctg tgt 518
Trp Thr Val Ala Arg Leu Tyr His Leu Leu Ala Glu Glu Asn Leu Cys

95

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ccg gcc tct aca agg gac atg gct tac cag gtg gcc ctt cgt gac ttt 566

Pro Ala Ser Thr Arg Asp Met Ala Tyr Gln Val Ala Leu Arg Asp Phe

110 115 120 125

gcc tcc cag ggt gac cac cag ctg ggc caa ctc cag aat gag gcc tgg 614

Ala Ser Gln Gly Asp His Gln Leu Gly Gln Leu Gln Asn Glu Ala Trp

130 135 140

gat cgg tgc agt tca gat atc aag ggg gac ccc agt ggt ttc cag cca 662

Asp Arg Cys Ser Ser Asp Ile Lys Gly Asp Pro Ser Gly Phe Gln Pro

145 150 155

ctc cat tct cat cag ggt tcc ctg cag cca cct tca gca tcc cct gca 710

Leu His Ser His Gln Gly Ser Leu Gln Pro Pro Ser Ala Ser Pro Ala

160 165 170

gtg acc aga agc cag cct cgt ccc att gac aca cca gac tgg agt tgg 758

Val Thr Arg Ser Gln Pro Arg Pro Ile Asp Thr Pro Asp Trp Ser Trp

175 180 185

gga cat acg tta cac tcc acc aac agc act gcc tca ctg gcc agc cac 806

Gly His Thr Leu His Ser Thr Asn Ser Thr Ala Ser Leu Ala Ser His

190 195 200 205

cta gag atc agc cag tca ccc act ctt gcc ttt ctc tct tca cac cat 854

Leu Glu Ile Ser Gln Ser Pro Thr Leu Ala Phe Leu Ser Ser His His

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gga acc cat ggg ccc agc aag cta tgt aac aca ccg ctg gac act cag 902

Gly Thr His Gly Pro Ser Lys Leu Cys Asn Thr Pro Leu Asp Thr Gln

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Trp Pro Pro Ser Val Glu Thr Ser Val Ser Leu Gly Leu Pro His Glu

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Ile Ser Val Pro Glu Val Ser Pro Glu Glu Ala Ser Pro Ile Leu Pro

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gac gcc ctg gct gct cca gac aca agt gtc cac tgt ccc att gaa tgc 1094

Asp Ala Leu Ala Ala Pro Asp Thr Ser Val His Cys Pro Ile Glu Cys

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aca gag ttg tct aca aac tcc agg tct ccc ctg acg tcc acc aca gaa 1142

Thr Glu Leu Ser Thr Asn Ser Arg Ser Pro Leu Thr Ser Thr Thr Glu

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agt gtt gga aag cag tgg cct att aca agt cag agg tca cct cag gtt 1190

Ser Val Gly Lys Gln Trp Pro Ile Thr Ser Gln Arg Ser Pro Gln Val

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cct gta gga gat gat tct ctg cag aac acc acg tca tcc agc cct cct 1238
 Pro Val Gly Asp Asp Ser Leu Gln Asn Thr Thr Ser Ser Ser Pro Pro
 335 340 345

gcc cag cca cca tcc ctc caa gcc tcc cct aag ctg cct cct tcc cct 1286
 Ala Gln Pro Pro Ser Leu Gln Ala Ser Pro Lys Leu Pro Pro Ser Pro
 350 355 360 365

ctg tcc tct gct tcc tcc ccg agc agc tac cct gct cct cca acc tcc 1334
 Leu Ser Ser Ala Ser Ser Pro Ser Ser Tyr Pro Ala Pro Pro Thr Ser
 370 375 380

aca tcc cct gtt ttg gac cac tca gaa aca tct gat cag aaa ttc tat 1382
 Thr Ser Pro Val Leu Asp His Ser Glu Thr Ser Asp Gln Lys Phe Tyr
 385 390 395

aac ttt gtg gtt atc cat gcc agg gct gat gaa cag gtg gcc cta cgt 1430
 Asn Phe Val Val Ile His Ala Arg Ala Asp Glu Gln Val Ala Leu Arg
 400 405 410

att cgg gag aag ctg gag acc ctc ggg gta cct gac ggg gcc acc ttc 1478
 Ile Arg Glu Lys Leu Glu Thr Leu Gly Val Pro Asp Gly Ala Thr Phe
 415 420 425

tgt gag gaa ttt cag gtg ccc ggg cgt ggt gag ctg cac tgt ctc caa 1526
 Cys Glu Glu Phe Gln Val Pro Gly Arg Gly Glu Leu His Cys Leu Gln
 430 435 440 445

gat gcc atc gat cac tcg ggg ttc acg atc ctg ctc ctg act gct agc 1574

Asp Ala Ile Asp His Ser Gly Phe Thr Ile Leu Leu Leu Thr Ala Ser

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455

460

ttt gat tgc agc ctg agc ctg cat caa atc aac cat gct ctc atg aac 1622

Phe Asp Cys Ser Leu Ser Leu His Gln Ile Asn His Ala Leu Met Asn

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470

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agc ctt aca cag tct ggg agg cag gac tgt gtg atc ccc ctc ctc cca 1670

Ser Leu Thr Gln Ser Gly Arg Gln Asp Cys Val Ile Pro Leu Leu Pro

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485

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ctt gag tgt tcc cag gcc cag ctc agc cca gat aca acc aga ctg ctc 1718

Leu Glu Cys Ser Gln Ala Gln Leu Ser Pro Asp Thr Thr Arg Leu Leu

495

500

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cac agc att gtg tgg ctg gat gaa cac tcc cca atc ttc gcc aga aag 1766

His Ser Ile Val Trp Leu Asp Glu His Ser Pro Ile Phe Ala Arg Lys

510

515

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525

gtg gca aac acc ttc aag aca cag aag ctc cag gca cag cgg gta cgc 1814

Val Ala Asn Thr Phe Lys Thr Gln Lys Leu Gln Ala Gln Arg Val Arg

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tgg aag aaa gcg cag gag gcc aga acc ctc aag gag cag agc ata cag 1862

Trp Lys Lys Ala Gln Glu Ala Arg Thr Leu Lys Glu Gln Ser Ile Gln

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Leu Glu Ala Glu Arg Gln Asn Val Ala Ala Ile Ser Ala Ala Tyr Thr

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Ala Tyr Val His Ser Tyr Arg Ala Trp Gln Ala Glu Met Asn Lys Leu			
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ggg gtg gct ttt ggg aag aac ttg tca ctg ggg act cca aca ccc agc			2006
Gly Val Ala Phe Gly Lys Asn Leu Ser Leu Gly Thr Pro Thr Pro Ser			
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tgg ccc gga tgt cca cag cca ata cct tct cat cct cag ggt ggt act			2054
Trp Pro Gly Cys Pro Gln Pro Ile Pro Ser His Pro Gln Gly Gly Thr			
610	615	620	
cca gtt ttc ccc tat tcc cca cag cct cca tcc ttc cct cag cct cca			2102
Pro Val Phe Pro Tyr Ser Pro Gln Pro Pro Ser Phe Pro Gln Pro Pro			
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tgc ttc cct cag cct cca tcc ttc cct cag cct cca tcc ttc cca ctg			2150
Cys Phe Pro Gln Pro Pro Ser Phe Pro Gln Pro Pro Ser Phe Pro Leu			
640	645	650	
cct cca gtc tct tcc cca cag tcc caa tcc ttt cca tca gcc tcc tcc			2198
Pro Pro Val Ser Ser Pro Gln Ser Gln Ser Phe Pro Ser Ala Ser Ser			
655	660	665	
cca gcc cca cag act cca gga cct cag cct ctc att att cac cat gcc			2246
Pro Ala Pro Gln Thr Pro Gly Pro Gln Pro Leu Ile Ile His His Ala			
670	675	680	685

cag atg gtt cag ctg ggt gtc aac aat cac atg tgg ggc cac aca ggg 2294

Gln Met Val Gln Leu Gly Val Asn Asn His Met Trp Gly His Thr Gly

690

695

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gcc cag tca tct gat gac aag act gag tgt tcg gag aac ccc tgt atg 2342

Ala Gln Ser Ser Asp Asp Lys Thr Glu Cys Ser Glu Asn Pro Cys Met

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ggc cct ctg act gat cag ggc gaa ccc ctt ctt gag act cca gag 2387

Gly Pro Leu Thr Asp Gln Gly Glu Pro Leu Leu Glu Thr Pro Glu

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2815

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<213> Homo sapiens

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30

Leu Cys Ser Gly Ser Gln Glu Ser Lys Leu Leu His Ala Met Val Leu

35

40

45

Leu Ala Leu Gly Gln Asp Thr Glu Ala Arg Val Ser Leu Glu Ser Leu

50

55

60

Lys Met Asn Thr Val Ala Gln Leu Val Ala His Gln Trp Ala Asp Met

65

70

75

80

Glu Thr Thr Glu Gly Pro Glu Glu Pro Pro Asp Leu Ser Trp Thr Val

85

90

95

Ala Arg Leu Tyr His Leu Leu Ala Glu Glu Asn Leu Cys Pro Ala Ser

100

105

110

Thr Arg Asp Met Ala Tyr Gln Val Ala Leu Arg Asp Phe Ala Ser Gln

115

120

125

Gly Asp His Gln Leu Gly Gln Leu Gln Asn Glu Ala Trp Asp Arg Cys

130

135

140

Ser Ser Asp Ile Lys Gly Asp Pro Ser Gly Phe Gln Pro Leu His Ser

145

150

155

160

His Gln Gly Ser Leu Gln Pro Pro Ser Ala Ser Pro Ala Val Thr Arg

165

170

175

Ser Gln Pro Arg Pro Ile Asp Thr Pro Asp Trp Ser Trp Gly His Thr

180

185

190

Leu His Ser Thr Asn Ser Thr Ala Ser Leu Ala Ser His Leu Glu Ile

195

200

205

Ser Gln Ser Pro Thr Leu Ala Phe Leu Ser Ser His His Gly Thr His

210

215

220

Gly Pro Ser Lys Leu Cys Asn Thr Pro Leu Asp Thr Gln Glu Pro Gln

225

230

235

240

Leu Val Pro Glu Gly Cys Gln Glu Pro Glu Glu Ile Ser Trp Pro Pro

245

250

255

Ser Val Glu Thr Ser Val Ser Leu Gly Leu Pro His Glu Ile Ser Val

260

265

270

Pro Glu Val Ser Pro Glu Glu Ala Ser Pro Ile Leu Pro Asp Ala Leu

275

280

285

Ala Ala Pro Asp Thr Ser Val His Cys Pro Ile Glu Cys Thr Glu Leu

290

295

300

Ser Thr Asn Ser Arg Ser Pro Leu Thr Ser Thr Thr Glu Ser Val Gly

305

310

315

320

Lys Gln Trp Pro Ile Thr Ser Gln Arg Ser Pro Gln Val Pro Val Gly

325

330

335

Asp Asp Ser Leu Gln Asn Thr Thr Ser Ser Ser Pro Pro Ala Gln Pro

340

345

350

Pro Ser Leu Gln Ala Ser Pro Lys Leu Pro Pro Ser Pro Leu Ser Ser

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365

Ala Ser Ser Pro Ser Ser Tyr Pro Ala Pro Pro Thr Ser Thr Ser Pro

370

375

380

Val Leu Asp His Ser Glu Thr Ser Asp Gln Lys Phe Tyr Asn Phe Val

385

390

395

400

Val Ile His Ala Arg Ala Asp Glu Gln Val Ala Leu Arg Ile Arg Glu

405

410

415

Lys Leu Glu Thr Leu Gly Val Pro Asp Gly Ala Thr Phe Cys Glu Glu

420

425

430

Phe Gln Val Pro Gly Arg Gly Glu Leu His Cys Leu Gln Asp Ala Ile

435 440 445

Asp His Ser Gly Phe Thr Ile Leu Leu Leu Thr Ala Ser Phe Asp Cys

450 455 460

Ser Leu Ser Leu His Gln Ile Asn His Ala Leu Met Asn Ser Leu Thr

465 470 475 480

Gln Ser Gly Arg Gln Asp Cys Val Ile Pro Leu Leu Pro Leu Glu Cys

485 490 495

Ser Gln Ala Gln Leu Ser Pro Asp Thr Thr Arg Leu Leu His Ser Ile

500 505 510

Val Trp Leu Asp Glu His Ser Pro Ile Phe Ala Arg Lys Val Ala Asn

515 520 525

Thr Phe Lys Thr Gln Lys Leu Gln Ala Gln Arg Val Arg Trp Lys Lys

530 535 540

Ala Gln Glu Ala Arg Thr Leu Lys Glu Gln Ser Ile Gln Leu Glu Ala

545 550 555 560

Glu Arg Gln Asn Val Ala Ala Ile Ser Ala Ala Tyr Thr Ala Tyr Val

565 570 575

His Ser Tyr Arg Ala Trp Gln Ala Glu Met Asn Lys Leu Gly Val Ala

580

585

590

Phe Gly Lys Asn Leu Ser Leu Gly Thr Pro Thr Pro Ser Trp Pro Gly

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600

605

Cys Pro Gln Pro Ile Pro Ser His Pro Gln Gly Gly Thr Pro Val Phe

610

615

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Pro Tyr Ser Pro Gln Pro Pro Ser Phe Pro Gln Pro Pro Cys Phe Pro

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Gln Pro Pro Ser Phe Pro Gln Pro Pro Ser Phe Pro Leu Pro Pro Val

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Ser Ser Pro Gln Ser Gln Ser Phe Pro Ser Ala Ser Ser Pro Ala Pro

660

665

670

Gln Thr Pro Gly Pro Gln Pro Leu Ile Ile His His Ala Gln Met Val

675

680

685

Gln Leu Gly Val Asn Asn His Met Trp Gly His Thr Gly Ala Gln Ser

690

695

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Ser Asp Asp Lys Thr Glu Cys Ser Glu Asn Pro Cys Met Gly Pro Leu

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Met Ala Cys Thr Gly Pro

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tca ctt cct agc gcc ttc gac att cta ggt gca gca ggc cag gac aag 162

Ser Leu Pro Ser Ala Phe Asp Ile Leu Gly Ala Ala Gly Gln Asp Lys

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ctc ttg tat ctg aag cac aaa ctg aag acc cca cgc cca ggc tgc cag 210

Leu Leu Tyr Leu Lys His Lys Leu Lys Thr Pro Arg Pro Gly Cys Gln

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ggg cag gac ctc ctg cat gcc atg gtt ctc ctg aag ctg ggc cag gaa 258

Gly Gln Asp Leu Leu His Ala Met Val Leu Leu Lys Leu Gly Gln Glu

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act gag gcc agg atc tct cta gag gca ttg aag gcc gat gcg gtg gcc 306
 Thr Glu Ala Arg Ile Ser Leu Glu Ala Leu Lys Ala Asp Ala Val Ala
 55 60 65 70

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 Arg Leu Val Ala Arg Gln Trp Ala Gly Val Asp Ser Thr Glu Asp Pro
 75 80 85

gag gag ccc cca gat gtg tcc tgg gct gtg gcc cgc ttg tac cac ctg 402
 Glu Glu Pro Pro Asp Val Ser Trp Ala Val Ala Arg Leu Tyr His Leu
 90 95 100

ctg gct gag gag aag ctg tgc ccc gcc tcg ctg cgg gac gtg gcc tac 450
 Leu Ala Glu Glu Lys Leu Cys Pro Ala Ser Leu Arg Asp Val Ala Tyr
 105 110 115

cag gaa gcc gtc cgc acc ctc agc tcc agg gac gac cac cgg ctg ggg 498
 Gln Glu Ala Val Arg Thr Leu Ser Ser Arg Asp Asp His Arg Leu Gly
 120 125 130

gaa ctt cag gat gag gcc cga aac cgg tgt ggg tgg gac att gct ggg 546
 Glu Leu Gln Asp Glu Ala Arg Asn Arg Cys Gly Trp Asp Ile Ala Gly
 135 140 145 150

gat cca ggg agc atc cgg acg ctc cag tcc aat ctg ggc tgc ctc cca 594
 Asp Pro Gly Ser Ile Arg Thr Leu Gln Ser Asn Leu Gly Cys Leu Pro
 155 160 165

cca tcc tcg gct ttg ccc tct ggg acc agg agc ctc cca cgc ccc att			642
Pro Ser Ser Ala Leu Pro Ser Gly Thr Arg Ser Leu Pro Arg Pro Ile			
170	175	180	
gac ggt gtt tcg gac tgg agc caa ggg tgc tcc ctg cga tcc act ggc			690
Asp Gly Val Ser Asp Trp Ser Gln Gly Cys Ser Leu Arg Ser Thr Gly			
185	190	195	
agc cct gcc tcc ctg gcc agc aac ttg gaa atc agc cag tcc cct acc			738
Ser Pro Ala Ser Leu Ala Ser Asn Leu Glu Ile Ser Gln Ser Pro Thr			
200	205	210	
atg ccc ttc ctc agc ctg cac cgc agc cca cat ggg ccc agc aag ctc			786
Met Pro Phe Leu Ser Leu His Arg Ser Pro His Gly Pro Ser Lys Leu			
215	220	225	230
tgt gac gac ccc cag gcc agc ttg gtg ccc gag cct gtc ccc ggt ggc			834
Cys Asp Asp Pro Gln Ala Ser Leu Val Pro Glu Pro Val Pro Gly Gly			
235	240	245	
tgc cag gag cct gag gag atg agc tgg ccg cca tcg ggg gag att gcc			882
Cys Gln Glu Pro Glu Glu Met Ser Trp Pro Pro Ser Gly Glu Ile Ala			
250	255	260	
agc cca cca gag ctg cca agc agc cca cct cct ggg ctt ccc gaa gtg			930
Ser Pro Pro Glu Leu Pro Ser Ser Pro Pro Pro Gly Leu Pro Glu Val			
265	270	275	
gcc cca gat gca acc tcc act ggc ctc cct gat acc ccc gca gct cca			978

Ala Pro Asp Ala Thr Ser Thr Gly Leu Pro Asp Thr Pro Ala Ala Pro

280

285

290

gaa acc agc acc aac tac cca gtg gag tgc acc gag ggg tct gca ggc 1026

Glu Thr Ser Thr Asn Tyr Pro Val Glu Cys Thr Glu Gly Ser Ala Gly

295

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310

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Pro Gln Ser Leu Pro Leu Pro Ile Leu Glu Pro Val Lys Asn Pro Cys

315

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Ser Val Lys Asp Gln Thr Pro Leu Gln Leu Ser Val Glu Asp Thr Thr

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tct cca aat acc aag ccg tgc cca cct act ccc acc acc cca gaa acá 1170

Ser Pro Asn Thr Lys Pro Cys Pro Pro Thr Pro Thr Pro Glu Thr

345

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tcc cct cct cct cct cct cct tca tct act cct tgt tca gct 1218

Ser Pro Pro Pro Pro Pro Pro Ser Ser Thr Pro Cys Ser Ala

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cac ctg acc ccc tcc tcc ctg ttc cct tcc tcc ctg gaa tca tca tcg 1266

His Leu Thr Pro Ser Ser Leu Phe Pro Ser Ser Leu Glu Ser Ser Ser

375

380

385

390

gaa cag aaa ttc tat aac ttt gtg atc ctc cac gcc agg gca gac gaa 1314

Glu Gln Lys Phe Tyr Asn Phe Val Ile Leu His Ala Arg Ala Asp Glu

395	400	405	
cac atc gcc ctg cggtt cgg gag aag ctg gag gcc ctt ggc gtg ccc 1362			
His Ile Ala Leu Arg Val Arg Glu Lys Leu Glu Ala Leu Gly Val Pro			
410	415	420	
gac ggg gcc acc ttc tgc gag gat ttc cag gtg ccg ggg cgc ggg gag 1410			
Asp Gly Ala Thr Phe Cys Glu Asp Phe Gln Val Pro Gly Arg Gly Glu			
425	430	435	
ctg agc tgc ctg cag gac gcc ata gac cac tca gct ttc atc atc cta 1458			
Leu Ser Cys Leu Gln Asp Ala Ile Asp His Ser Ala Phe Ile Ile Leu			
440	445	450	
ctt ctc acc tcc aac ttc gac tgt cgc ctg agc ctg cac cag gtg aac 1506			
Leu Leu Thr Ser Asn Phe Asp Cys Arg Leu Ser Leu His Gln Val Asn			
455	460	465	470
caa gcc atg atg agc aac ctc acg cga cag ggg tcg cca gac tgt gtc 1554			
Gln Ala Met Met Ser Asn Leu Thr Arg Gln Gly Ser Pro Asp Cys Val			
475	480	485	
atc ccc ttc ctg ccc ctg gag agc tcc ccg gcc cag ctc agc tcc gac 1602			
Ile Pro Phe Leu Pro Leu Glu Ser Ser Pro Ala Gln Leu Ser Ser Asp			
490	495	500	
acg gcc agc ctg ctc tcc ggg ctg gtg cgg ctg gac gaa cac tcc cag 1650			
Thr Ala Ser Leu Leu Ser Gly Leu Val Arg Leu Asp Glu His Ser Gln			
505	510	515	

atc ttc gcc agg aag gtg gcc aac acc ttc aag ccc cac agg ctt cag 1698

Ile Phe Ala Arg Lys Val Ala Asn Thr Phe Lys Pro His Arg Leu Gln

520 525 530

gcc cga aag gcc atg tgg agg aag gaa cag gac acc cga gcc ctg cgg 1746

Ala Arg Lys Ala Met Trp Arg Lys Glu Gln Asp Thr Arg Ala Leu Arg

535 540 545 550

gaa cag agc caa cac ctg gac ggt gag cgg atg cag gcg gcg gca ctg 1794

Glu Gln Ser Gln His Leu Asp Gly Glu Arg Met Gln Ala Ala Ala Leu

555 560 565

aac gca gcc tac tca gcc tac ctc cag agc tac ttg tcc tac cag gca 1842

Asn Ala Ala Tyr Ser Ala Tyr Leu Gln Ser Tyr Leu Ser Tyr Gln Ala

570 575 580

cag atg gag cag ctc cag gtg gct ttt ggg agc cac atg tca ttt ggg 1890

Gln Met Glu Gln Leu Gln Val Ala Phe Gly Ser His Met Ser Phe Gly

585 590 595

act ggg gcg ccc tat ggg gct cga atg ccc ttt ggg ggc cag gtg ccc 1938

Thr Gly Ala Pro Tyr Gly Ala Arg Met Pro Phe Gly Gly Gln Val Pro

600 605 610

ctg gga gcc ccg cca ccc ttt ccc act tgg ccg ggg tgc ccg cag ccg 1986

Leu Gly Ala Pro Pro Pro Phe Pro Thr Trp Pro Gly Cys Pro Gln Pro

615 620 625 630

cca ccc ctg cac gca tgg cag gct ggc acc ccc cca ccg ccc tcc cca 2034
 Pro Pro Leu His Ala Trp Gln Ala Gly Thr Pro Pro Pro Pro Ser Pro
 635 640 645

cag cca gca gcc ttt cca cag tca ctg ccc ttc ccg cag tcc cca gcc 2082
 Gln Pro Ala Ala Phe Pro Gln Ser Leu Pro Phe Pro Gln Ser Pro Ala
 650 655 660

ttc cct acg gcc tca ccc gca ccc cct cag agc cca ggg ctg caa ccc 2130
 Phe Pro Thr Ala Ser Pro Ala Pro Pro Gln Ser Pro Gly Leu Gln Pro
 665 670 675

ctc att atc cac cac gca cag atg gta cag ctg ggg ctg aac aac cac 2178
 Leu Ile Ile His His Ala Gln Met Val Gln Leu Gly Leu Asn Asn His
 680 685 690

atg tgg aac cag aga ggg tcc cag gcg ccc gag gac aag acg cag gag 2226
 Met Trp Asn Gln Arg Gly Ser Gln Ala Pro Glu Asp Lys Thr Gln Glu
 695 700 705 710

gca gaa tgaccgcgtg tccttgcctg accacacctggg gaacacccct ggacccaggc 2282
 Ala Glu

atcggccagg accccataga gcaccccggt ctgccctgtg ccctgtggac agtggaaagat 2342

gaggcatct gccacttca ggacattgtc cgggagccct tcatttagga caaaacggc 2402

gcgatgatgc cctggcttc agggtggta gaactggata cggtgtttac aattccaatc 2462

tctctatttc tgggtgaagg gtcttggtgg tgggggtatt gctacggtct tttaattata 2522

ataaatattt attgaatgct tc 2544

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<212> PRT

<213> Homo sapiens

<400> 154

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1

5

10

15

Ala Ala Gly Gln Asp Lys Leu Leu Tyr Leu Lys His Lys Leu Lys Thr

20

25

30

Pro Arg Pro Gly Cys Gln Gly Gln Asp Leu Leu His Ala Met Val Leu

35

40

45

Leu Lys Leu Gly Gln Glu Thr Glu Ala Arg Ile Ser Leu Glu Ala Leu

50

55

60

Lys Ala Asp Ala Val Ala Arg Leu Val Ala Arg Gln Trp Ala Gly Val

65

70

75

80

Asp Ser Thr Glu Asp Pro Glu Glu Pro Pro Asp Val Ser Trp Ala Val

85

90

95

Ala Arg Leu Tyr His Leu Leu Ala Glu Glu Lys Leu Cys Pro Ala Ser

100 105 110

Leu Arg Asp Val Ala Tyr Gln Glu Ala Val Arg Thr Leu Ser Ser Arg

115 120 125

Asp Asp His Arg Leu Gly Glu Leu Gln Asp Glu Ala Arg Asn Arg Cys

130 135 140

Gly Trp Asp Ile Ala Gly Asp Pro Gly Ser Ile Arg Thr Leu Gln Ser

145 150 155 160

Asn Leu Gly Cys Leu Pro Pro Ser Ser Ala Leu Pro Ser Gly Thr Arg

165 170 175

Ser Leu Pro Arg Pro Ile Asp Gly Val Ser Asp Trp Ser Gln Gly Cys

180 185 190

Ser Leu Arg Ser Thr Gly Ser Pro Ala Ser Leu Ala Ser Asn Leu Glu

195 200 205

Ile Ser Gln Ser Pro Thr Met Pro Phe Leu Ser Leu His Arg Ser Pro

210 215 220

His Gly Pro Ser Lys Leu Cys Asp Asp Pro Gln Ala Ser Leu Val Pro

225 230 235 240

Glu Pro Val Pro Gly Gly Cys Gln Glu Pro Glu Glu Met Ser Trp Pro

245 250 255

Pro Ser Gly Glu Ile Ala Ser Pro Pro Glu Leu Pro Ser Ser Pro Pro

260 265 270

Pro Gly Leu Pro Glu Val Ala Pro Asp Ala Thr Ser Thr Gly Leu Pro

275 280 285

Asp Thr Pro Ala Ala Pro Glu Thr Ser Thr Asn Tyr Pro Val Glu Cys

290 295 300

Thr Glu Gly Ser Ala Gly Pro Gln Ser Leu Pro Leu Pro Ile Leu Glu

305 310 315 320

Pro Val Lys Asn Pro Cys Ser Val Lys Asp Gln Thr Pro Leu Gln Leu

325 330 335

Ser Val Glu Asp Thr Thr Ser Pro Asn Thr Lys Pro Cys Pro Pro Thr

340 345 350

Pro Thr Thr Pro Glu Thr Ser Pro Pro Pro Pro Pro Pro Pro Ser

355 360 365

Ser Thr Pro Cys Ser Ala His Leu Thr Pro Ser Ser Leu Phe Pro Ser

370 375 380

Ser Leu Glu Ser Ser Ser Glu Gln Lys Phe Tyr Asn Phe Val Ile Leu

385 390 395 400

His Ala Arg Ala Asp Glu His Ile Ala Leu Arg Val Arg Glu Lys Leu

405	410	415
Glu Ala Leu Gly Val Pro Asp Gly Ala Thr Phe Cys Glu Asp Phe Gln		
420	425	430
Val Pro Gly Arg Gly Glu Leu Ser Cys Leu Gln Asp Ala Ile Asp His		
435	440	445
Ser Ala Phe Ile Ile Leu Leu Leu Thr Ser Asn Phe Asp Cys Arg Leu		
450	455	460
Ser Leu His Gln Val Asn Gln Ala Met Met Ser Asn Leu Thr Arg Gln		
465	470	475
480		
Gly Ser Pro Asp Cys Val Ile Pro Phe Leu Pro Leu Glu Ser Ser Pro		
485	490	495
Ala Gln Leu Ser Ser Asp Thr Ala Ser Leu Leu Ser Gly Leu Val Arg		
500	505	510
Leu Asp Glu His Ser Gln Ile Phe Ala Arg Lys Val Ala Asn Thr Phe		
515	520	525
Lys Pro His Arg Leu Gln Ala Arg Lys Ala Met Trp Arg Lys Glu Gln		
530	535	540
Asp Thr Arg Ala Leu Arg Glu Gln Ser Gln His Leu Asp Gly Glu Arg		
545	550	555
560		

Met Gln Ala Ala Ala Leu Asn Ala Ala Tyr Ser Ala Tyr Leu Gln Ser

565 570 575

Tyr Leu Ser Tyr Gln Ala Gln Met Glu Gln Leu Gln Val Ala Phe Gly

580 585 590

Ser His Met Ser Phe Gly Thr Gly Ala Pro Tyr Gly Ala Arg Met Pro

595 600 605

Phe Gly Gly Gln Val Pro Leu Gly Ala Pro Pro Pro Phe Pro Thr Trp

610 615 620

Pro Gly Cys Pro Gln Pro Pro Pro Leu His Ala Trp Gln Ala Gly Thr

625 630 635 640

Pro Pro Pro Pro Ser Pro Gln Pro Ala Ala Phe Pro Gln Ser Leu Pro

645 650 655

Phe Pro Gln Ser Pro Ala Phe Pro Thr Ala Ser Pro Ala Pro Pro Gln

660 665 670

Ser Pro Gly Leu Gln Pro Leu Ile Ile His His Ala Gln Met Val Gln

675 680 685

Leu Gly Leu Asn Asn His Met Trp Asn Gln Arg Gly Ser Gln Ala Pro

690 695 700

Glu Asp Lys Thr Gln Glu Ala Glu

705 710

<210> 155

<211> 3456

<212> DNA

<213> Homo sapiens

<220>

<221> CDS

<222> (281)..(3016)

<400> 155

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ggaggagagc tctgtggatg gcaaagggga ccggaagagc acaggcctga aactctccaa 180

gaagaaaagca aggaggagac acacggatga cccaaagcaag gaatgcttca ctctgaaatt 240

tgacctgaat gtggacattg agacagagat cgtcccagcc atg aag aag aag tca 295

Met Lys Lys Lys Ser

1

5

ctg ggg gag gtg ctg cct gta ttt gaa agg aag ggc att gcg ctg 343

Leu Gly Glu Val Leu Leu Pro Val Phe Glu Arg Lys Gly Ile Ala Leu

10

15

20

ggc aaa gtg gac atc tac ctg gac cag tcc aac aca ccc ctg tcc ctc			391
Gly Lys Val Asp Ile Tyr Leu Asp Gln Ser Asn Thr Pro Leu Ser Leu			
25	30	35	
acc ttc gag gcc tac agg ttc ggg gga cac tac ctt cgt gtc aaa gcc			439
Thr Phe Glu Ala Tyr Arg Phe Gly Gly His Tyr Leu Arg Val Lys Ala			
40	45	50	
cca gcc aag cct gga gat gag ggc aag gtg gag cag ggc atg aag gac			487
Pro Ala Lys Pro Gly Asp Glu Gly Lys Val Glu Gln Gly Met Lys Asp			
55	60	65	
tcc aag tcc ctg agt ttg ccg att ctg cgg cca gct ggg acc ggg ccc			535
Ser Lys Ser Leu Ser Leu Pro Ile Leu Arg Pro Ala Gly Thr Gly Pro			
70	75	80	85
ccc gcc ctg gag cgt gtg gac gcc cag agc cgc cgg gag agc ctg gac			583
Pro Ala Leu Glu Arg Val Asp Ala Gln Ser Arg Arg Glu Ser Leu Asp			
90	95	100	
atc ttg gcc cct ggc cgc cgc aag aac atg tcg gag ttc ctg ggg			631
Ile Leu Ala Pro Gly Arg Arg Lys Asn Met Ser Glu Phe Leu Gly			
105	110	115	
gag gcg agc atc ccc ggg cag gag ccc acg ccc tcc agc tgc tct			679
Glu Ala Ser Ile Pro Gly Gln Glu Pro Pro Thr Pro Ser Ser Cys Ser			
120	125	130	
ctg ccc agc ggc agc agt ggc agc acc aac act ggc gac agc tgg aag			727

Leu Pro Ser Gly Ser Ser Gly Ser Thr Asn Thr Gly Asp Ser Trp Lys

135

140

145

aac cgg gcg gcc agt cgc ttc agc ggc ttt ttc agc tcc ggc ccc agc 775

Asn Arg Ala Ala Ser Arg Phe Ser Gly Phe Phe Ser Ser Gly Pro Ser

150

155

160

165

acc agc gcc ttt ggc cgg gag gta gac aag atg gag cag ctg gag ggc 823

Thr Ser Ala Phe Gly Arg Glu Val Asp Lys Met Glu Gln Leu Glu Gly

170

175

180

aag ctg cac acc tac agc ctc ttc ggg ctg ccc agg ctg ccc cgg ggg 871

Lys Leu His Thr Tyr Ser Leu Phe Gly Leu Pro Arg Leu Pro Arg Gly

185

190

195

ctg cgc ttc gac cat gac tcc tgg gag gag gag tac gat gaa gac gag 919

Leu Arg Phe Asp His Asp Ser Trp Glu Glu Glu Tyr Asp Glu Asp Glu

200

205

210

gat gag gac aat gcc tgc ctg agg ctg gag gac agc tgg cgg gag ctc 967

Asp Glu Asp Asn Ala Cys Leu Arg Leu Glu Asp Ser Trp Arg Glu Leu

215

220

225

att gat ggg cat gag aag ctg acc cgg cgg cag tgc cac cag cag gag 1015

Ile Asp Gly His Glu Lys Leu Thr Arg Arg Gln Cys His Gln Gln Glu

230

235

240

245

gcg gtg tgg gag ctg ctg cac acg gag gcc tcc tac atc agg aaa ctg 1063

Ala Val Trp Glu Leu Leu His Thr Glu Ala Ser Tyr Ile Arg Lys Leu

250	255	260	
cggtgtatc atc aac ctg ttc ttgc tgc ctc ctg aac ctg caa gag 1111 Arg Val Ile Ile Asn Leu Phe Leu Cys Cys Leu Leu Asn Leu Gln Glu			
265	270	275	
tcaggctgctgttgt gag gtggag gcg gag cgc ctgttc agcaacatc 1159 Ser Gly Leu Leu Cys Glu Val Glu Ala Glu Arg Leu Phe Ser Asn Ile			
280	285	290	
ccggagatcgccg cag ctgcac cgc agg ctgtgg gct agc gtg atg gcg 1207 Pro Glu Ile Ala Gln Leu His Arg Arg Leu Trp Ala Ser Val Met Ala			
295	300	305	
ccggatgtctggag aag gcg cgg cgc acg cga gcg ctg cta cag ccc ggg 1255 Pro Val Leu Glu Lys Ala Arg Arg Thr Arg Ala Leu Leu Gln Pro Gly			
310	315	320	325
gac ttcctc aaa ggc ttc aag atg ttc ggc tcg ctc ttc aag ccc tac 1303 Asp Phe Leu Lys Gly Phe Lys Met Phe Gly Ser Leu Phe Lys Pro Tyr			
330	335	340	
atccgc tac tgc atg gag gag ggc tgc atg gag tac atg cgc ggc 1351 Ile Arg Tyr Cys Met Glu Glu Gly Cys Met Glu Tyr Met Arg Gly			
345	350	355	
ctgcctgcgac aac gac ctc ttc cgg gcc tac atc acg tgg gcg gag 1399 Leu Leu Arg Asp Asn Asp Leu Phe Arg Ala Tyr Ile Thr Trp Ala Glu			
360	365	370	

aag cac cca cag tgc cag agg ctg aag ctg agc gac atg ctg gcc aaa 1447
 Lys His Pro Gln Cys Gln Arg Leu Lys Leu Ser Asp Met Leu Ala Lys
 375 380 385

ccc cac cag cgg ctc acc aag tac ccg ctg ctc aag tcg gtg ctg 1495
 Pro His Gln Arg Leu Thr Lys Tyr Pro Leu Leu Lys Ser Val Leu
 390 395 400 405

agg aag acc gag gag ccg cgc gcc aag gag gcc gtc gtc gcc atg atc 1543
 Arg Lys Thr Glu Glu Pro Arg Ala Lys Glu Ala Val Val Ala Met Ile
 410 415 420

ggc tcc gtg gag cgc ttc atc cac cac gtg aac gcg tgc atg cgg cag 1591
 Gly Ser Val Glu Arg Phe Ile His His Val Asn Ala Cys Met Arg Gln
 425 430 435

cgg cag gag cgg cag cgg ctg gcg gcc gtg gtg agc cgc atc gac gcc 1639
 Arg Gln Glu Arg Gln Arg Leu Ala Ala Val Val Ser Arg Ile Asp Ala
 440 445 450

tac gag gtg gtg gaa agc agc gac gaa gtg gac aag ctc ctg aag 1687
 Tyr Glu Val Val Glu Ser Ser Asp Glu Val Asp Lys Leu Leu Lys
 455 460 465

gaa ttt ctg cac ctg gac ttg aca gcg ccc atc cct ggc gcc tcc cgg 1735
 Glu Phe Leu His Leu Asp Leu Thr Ala Pro Ile Pro Gly Ala Ser Pro
 470 475 480 485

gag gag acg cgg cag ctg ctg gag ggg agc ctg agg atg aag gag			1783
Glu Glu Thr Arg Gln Leu Leu Glu Gly Ser Leu Arg Met Lys Glu			
490	495	500	
ggg aag gac agc aag atg gat gtg tac tgc ttc ctc ttc acg gat ctg			1831
Gly Lys Asp Ser Lys Met Asp Val Tyr Cys Phe Leu Phe Thr Asp Leu			
505	510	515	
ctg ttg gtg acc aaa gca gtg aag aag gca gag agg acc agg gtc atc			1879
Leu Leu Val Thr Lys Ala Val Lys Lys Ala Glu Arg Thr Arg Val Ile			
520	525	530	
agg cca ccc ctg ctc gtg gac aag att gtg tgc cgg gag cta cgg gac			1927
Arg Pro Pro Leu Leu Val Asp Lys Ile Val Cys Arg Glu Leu Arg Asp			
535	540	545	
cct ggg tcc ttc ctc ctt atc tac ctg aat gag ttt cac agt gct gta			1975
Pro Gly Ser Phe Leu Leu Ile Tyr Leu Asn Glu Phe His Ser Ala Val			
550	555	560	565
ggg gcc tac acg ttc cag gcc agt ggc cag gcc ttg tgc cgt ggc tgg			2023
Gly Ala Tyr Thr Phe Gln Ala Ser Gly Gln Ala Leu Cys Arg Gly Trp			
570	575	580	
gtg gac acc att tac aat gcc cag aac cag ctg caa cag ctg cgt gca			2071
Val Asp Thr Ile Tyr Asn Ala Gln Asn Gln Leu Gln Gln Leu Arg Ala			
585	590	595	
cag gag ccc cca ggc agt cag cag ccc ctg cag agc ctg gaa gag gag			2119

Gln Glu Pro Pro Gly Ser Gln Gln Pro Leu Gln Ser Leu Glu Glu Glu

600

605

610

gag gat gag cag gag gag gaa gag gag gag gag gag gag gag gag gaa 2167

Glu Asp Glu Gln Glu Glu

615

620

625

ggc gag gac agt ggc act tca gct gcc agc tcc cct acc atc atg cgg 2215

Gly Glu Asp Ser Gly Thr Ser Ala Ala Ser Ser Pro Thr Ile Met Arg

630

635

640

645

aaa agc agc ggc agc ccc gac tct cag cac tgt gcc tca gat ggc tcc 2263

Lys Ser Ser Gly Ser Pro Asp Ser Gln His Cys Ala Ser Asp Gly Ser

650

655

660

acg gag acc ctg gcc atg gtt gtg gta gag cct ggg gac acg ctg tcc 2311

Thr Glu Thr Leu Ala Met Val Val Val Glu Pro Gly Asp Thr Leu Ser

665

670

675

tcc ccc gag ttc gac agc ggt cct ttc agc tcc cag tct gat gag acc 2359

Ser Pro Glu Phe Asp Ser Gly Pro Phe Ser Ser Gln Ser Asp Glu Thr

680

685

690

tct ctc agc acc act gcc tca tct gcc acg ccc acc agt gag ctg ctg 2407

Ser Leu Ser Thr Thr Ala Ser Ser Ala Thr Pro Thr Ser Glu Leu Leu

695

700

705

ccc ctg ggt ccg gtg gac ggc cgc tcc tgc tcc atg gac tct gcc tac 2455

Pro Leu Gly Pro Val Asp Gly Arg Ser Cys Ser Met Asp Ser Ala Tyr

710

715

720

725

ggc acc ctc tcc cca acc tcc tta caa gac ttt gtg gcc cca ggc cca 2503
 Gly Thr Leu Ser Pro Thr Ser Leu Gln Asp Phe Val Ala Pro Gly Pro

730

735

740

atg gca gag cta gtg cct cgg gcc cca gag tcc cca cga gtt cct tcc 2551
 Met Ala Glu Leu Val Pro Arg Ala Pro Glu Ser Pro Arg Val Pro Ser

745

750

755

cct cca ccc tcg ccc cgt ctc cgc cgc cgc acc cct gtc cag ctg ttg 2599
 Pro Pro Pro Ser Pro Arg Leu Arg Arg Arg Thr Pro Val Gln Leu Leu

760

765

770

agc tgc ccg ccc cac ctg ctc aag tct aag tcc gag gcc agc ctc ctc 2647
 Ser Cys Pro Pro His Leu Leu Lys Ser Lys Ser Glu Ala Ser Leu Leu

775

780

785

cag ctg ctg gca ggg gct ggc acc cat ggg aca ccc tct gcc ccc agc 2695
 Gln Leu Leu Ala Gly Ala Gly Thr His Gly Thr Pro Ser Ala Pro Ser

790 795 800 805

cgc agc ctg tca gag ctc tgc ctg gct gtt cca gcc cca ggt att agg 2743
 Arg Ser Leu Ser Glu Leu Cys Leu Ala Val Pro Ala Pro Gly Ile Arg

810

815

820

act cag ggc tcc cct cag gaa gct ggg ccc agc tgg gat tgc cga ggg 2791
 Thr Gln Gly Ser Pro Gln Glu Ala Gly Pro Ser Trp Asp Cys Arg Gly

825

830

835

gcc cct agc cct ggc agc ggt cct ggg cta gtc ggc tgc ctg gcc ggg 2839
 Ala Pro Ser Pro Gly Ser Gly Pro Gly Leu Val Gly Cys Leu Ala Gly

840 845 850

gaa cct gca ggc tcc cac agg aag agg tgt gga gac ctg ccc tcg ggg 2887
 Glu Pro Ala Gly Ser His Arg Lys Arg Cys Gly Asp Leu Pro Ser Gly

855 860 865

gcc tct ccc agg gtc cag cct gag ccc cca cca ggg gtc tct gcc cag 2935
 Ala Ser Pro Arg Val Gln Pro Glu Pro Pro Gly Val Ser Ala Gln
 870 875 880 885

cac agg aag ctg acc ctg gcc cag ctc tac cga atc agg acc acc ctg 2983
 His Arg Lys Leu Thr Leu Ala Gln Leu Tyr Arg Ile Arg Thr Thr Leu
 890 895 900

ctg ctt aac tcc acg ctc act gcc tcg gag gtc tgagcagagg gaggccccca 3036
 Leu Leu Asn Ser Thr Leu Thr Ala Ser Glu Val
 905 910

agagtccat tgaccaagag acagcagaca gcctgcctcc tggggcggtgc cggcacctgc 3096

ttagctact gcctccgtta tgcattgagcc ggatgctggg caggatccct gcctacgccc 3156

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ccccacacctgc ccaggcagcc cctcgtaacc tactccccga agttaccaggc tcagctcgag 3276

tcttcagggc tgggctccta ggctgccat cctacttcta ccctcaactgg cctccagtgg 3336

gattcactcc tgccctgccc ccaccttccc agtcccacag gccaccctg gcttgggctg 3396

ggttctgtga agttacgtat ttattgagct tttggttctt ttataaagac ttgtctagac 3456

<210> 156

<211> 912

<212> PRT

<213> Homo sapiens

<400> 156

Met Lys Lys Lys Ser Leu Gly Glu Val Leu Leu Pro Val Phe Glu Arg

1

5

10

15

Lys Gly Ile Ala Leu Gly Lys Val Asp Ile Tyr Leu Asp Gln Ser Asn

20

25

30

Thr Pro Leu Ser Leu Thr Phe Glu Ala Tyr Arg Phe Gly Gly His Tyr

35

40

45

Leu Arg Val Lys Ala Pro Ala Lys Pro Gly Asp Glu Gly Lys Val Glu

50

55

60

Gln Gly Met Lys Asp Ser Lys Ser Leu Ser Leu Pro Ile Leu Arg Pro

65

70

75

80

Ala Gly Thr Gly Pro Pro Ala Leu Glu Arg Val Asp Ala Gln Ser Arg

85	90	95
Arg Glu Ser Leu Asp Ile Leu Ala Pro Gly Arg Arg Lys Asn Met		
100	105	110
Ser Glu Phe Leu Gly Glu Ala Ser Ile Pro Gly Gln Glu Pro Pro Thr		
115	120	125
Pro Ser Ser Cys Ser Leu Pro Ser Gly Ser Ser Gly Ser Thr Asn Thr		
130	135	140
Gly Asp Ser Trp Lys Asn Arg Ala Ala Ser Arg Phe Ser Gly Phe Phe		
145	150	155
160		
Ser Ser Gly Pro Ser Thr Ser Ala Phe Gly Arg Glu Val Asp Lys Met		
165	170	175
Glu Gln Leu Glu Gly Lys Leu His Thr Tyr Ser Leu Phe Gly Leu Pro		
180	185	190
Arg Leu Pro Arg Gly Leu Arg Phe Asp His Asp Ser Trp Glu Glu Glu		
195	200	205
Tyr Asp Glu Asp Glu Asp Asn Ala Cys Leu Arg Leu Glu Asp		
210	215	220
Ser Trp Arg Glu Leu Ile Asp Gly His Glu Lys Leu Thr Arg Arg Gln		
225	230	235
240		

Cys His Gln Gln Glu Ala Val Trp Glu Leu Leu His Thr Glu Ala Ser

245

250

255

Tyr Ile Arg Lys Leu Arg Val Ile Ile Asn Leu Phe Leu Cys Cys Leu

260

265

270

Leu Asn Leu Gln Glu Ser Gly Leu Leu Cys Glu Val Glu Ala Glu Arg

275

280

285

Leu Phe Ser Asn Ile Pro Glu Ile Ala Gln Leu His Arg Arg Leu Trp

290

295

300

Ala Ser Val Met Ala Pro Val Leu Glu Lys Ala Arg Arg Thr Arg Ala

305

310

315

320

Leu Leu Gln Pro Gly Asp Phe Leu Lys Gly Phe Lys Met Phe Gly Ser

325

330

335

Leu Phe Lys Pro Tyr Ile Arg Tyr Cys Met Glu Glu Glu Gly Cys Met

340

345

350

Glu Tyr Met Arg Gly Leu Leu Arg Asp Asn Asp Leu Phe Arg Ala Tyr

355

360

365

Ile Thr Trp Ala Glu Lys His Pro Gln Cys Gln Arg Leu Lys Leu Ser

370

375

380

Asp Met Leu Ala Lys Pro His Gln Arg Leu Thr Lys Tyr Pro Leu Leu

385

390

395

400

Leu Lys Ser Val Leu Arg Lys Thr Glu Glu Pro Arg Ala Lys Glu Ala

405

410

415

Val Val Ala Met Ile Gly Ser Val Glu Arg Phe Ile His His Val Asn

420

425

430

Ala Cys Met Arg Gln Arg Gln Glu Arg Gln Arg Leu Ala Ala Val Val

435

440

445

Ser Arg Ile Asp Ala Tyr Glu Val Val Glu Ser Ser Ser Asp Glu Val

450

455

460

Asp Lys Leu Leu Lys Glu Phe Leu His Leu Asp Leu Thr Ala Pro Ile

465

470

475

480

Pro Gly Ala Ser Pro Glu Glu Thr Arg Gln Leu Leu Leu Glu Gly Ser

485

490

495

Leu Arg Met Lys Glu Gly Lys Asp Ser Lys Met Asp Val Tyr Cys Phe

500

505

510

Leu Phe Thr Asp Leu Leu Val Thr Lys Ala Val Lys Lys Ala Glu

515

520

525

Arg Thr Arg Val Ile Arg Pro Pro Leu Leu Val Asp Lys Ile Val Cys

530

535

540

Arg Glu Leu Arg Asp Pro Gly Ser Phe Leu Leu Ile Tyr Leu Asn Glu

545

550

555

560

Phe His Ser Ala Val Gly Ala Tyr Thr Phe Gln Ala Ser Gly Gln Ala

565

570

575

Leu Cys Arg Gly Trp Val Asp Thr Ile Tyr Asn Ala Gln Asn Gln Leu

580

585

590

Gln Gln Leu Arg Ala Gln Glu Pro Pro Gly Ser Gln Gln Pro Leu Gln

595

600

605

Ser Leu Glu Glu Glu Asp Glu Gln Glu Glu Glu Glu Glu Glu Glu

610

615

620

Glu Glu Glu Glu Glu Gly Glu Asp Ser Gly Thr Ser Ala Ala Ser Ser

625

630

635

640

Pro Thr Ile Met Arg Lys Ser Ser Gly Ser Pro Asp Ser Gln His Cys

645

650

655

Ala Ser Asp Gly Ser Thr Glu Thr Leu Ala Met Val Val Val Glu Pro

660

665

670

Gly Asp Thr Leu Ser Ser Pro Glu Phe Asp Ser Gly Pro Phe Ser Ser

675

680

685

Gln Ser Asp Glu Thr Ser Leu Ser Thr Thr Ala Ser Ser Ala Thr Pro

690

695

700

Thr Ser Glu Leu Leu Pro Leu Gly Pro Val Asp Gly Arg Ser Cys Ser
 705 710 715 720

Met Asp Ser Ala Tyr Gly Thr Leu Ser Pro Thr Ser Leu Gln Asp Phe
 725 730 735

Val Ala Pro Gly Pro Met Ala Glu Leu Val Pro Arg Ala Pro Glu Ser
 740 745 750

Pro Arg Val Pro Ser Pro Pro Pro Ser Pro Arg Leu Arg Arg Arg Thr
 755 760 765

Pro Val Gln Leu Leu Ser Cys Pro Pro His Leu Leu Lys Ser Lys Ser
 770 775 780

Glu Ala Ser Leu Leu Gln Leu Leu Ala Gly Ala Gly Thr His Gly Thr
 785 790 795 800

Pro Ser Ala Pro Ser Arg Ser Leu Ser Glu Leu Cys Leu Ala Val Pro
 805 810 815

Ala Pro Gly Ile Arg Thr Gln Gly Ser Pro Gln Glu Ala Gly Pro Ser
 820 825 830

Trp Asp Cys Arg Gly Ala Pro Ser Pro Gly Ser Gly Pro Gly Leu Val
 835 840 845

Gly Cys Leu Ala Gly Glu Pro Ala Gly Ser His Arg Lys Arg Cys Gly
 850 855 860

Asp Leu Pro Ser Gly Ala Ser Pro Arg Val Gln Pro Glu Pro Pro

865 870 875 880

Gly Val Ser Ala Gln His Arg Lys Leu Thr Leu Ala Gln Leu Tyr Arg

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cgaggcctgt gacagcaagt tccacagcac c atg cat tat gat ggg cat gtc 172

Met His Tyr Asp Gly His Val

1 5

cgc ttc gac ctt ccc cca caa ggc tct gtg ctg gcc cg_g aac gtg tcc 220

Arg Phe Asp Leu Pro Pro Gln Gly Ser Val Leu Ala Arg Asn Val Ser

10

15

20

acc cgg tca tgc ccg ccg cgc acc agc ccc gca gtg gac ttg gag gag 268

Thr Arg Ser Cys Pro Pro Arg Thr Ser Pro Ala Val Asp Leu Glu Glu

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gag gag gag gag agc tct gtg gat ggc aaa ggg gac cgg aag agc aca 316

Glu Glu Glu Glu Ser Ser Val Asp Gly Lys Gly Asp Arg Lys Ser Thr

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ggc ctg aaa ctc tcc aag aag aaa gca agg agg aga cac acg gat gac 364

Gly Leu Lys Leu Ser Lys Lys Ala Arg Arg Arg His Thr Asp Asp

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cca agc aag gaa tgc ttc act ctg aaa ttt gac ctg aat gtg gac att 412

Pro Ser Lys Glu Cys Phe Thr Leu Lys Phe Asp Leu Asn Val Asp Ile

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gag aca gag atc gtc cca gcc atg aag aag aag tca ctg ggg gag gtg 460

Glu Thr Glu Ile Val Pro Ala Met Lys Lys Ser Leu Gly Glu Val

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95

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ctg ctg cct gta ttt gaa agg aag ggc att gcg ctg ggc aaa gtg gac 508

Leu Leu Pro Val Phe Glu Arg Lys Gly Ile Ala Leu Gly Lys Val Asp

105

110

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atc tac ctg gac cag tcc aac aca ccc ctg tcc ctc acc ttc gag gcc	556		
Ile Tyr Leu Asp Gln Ser Asn Thr Pro Leu Ser Leu Thr Phe Glu Ala			
120	125	130	135
 tac agg ttc ggg gga cac tac ctt cgt gtc aaa gcc cca gcc aag cct	604		
Tyr Arg Phe Gly Gly His Tyr Leu Arg Val Lys Ala Pro Ala Lys Pro			
140	145	150	
 gga gat gag ggc aag gtg gag cag ggc atg aag gac tcc aag tcc ctg	652		
Gly Asp Glu Gly Lys Val Glu Gln Gly Met Lys Asp Ser Lys Ser Leu			
155	160	165	
 agt ttg ccg att ctg cgg cca gct ggg acc ggg ccc ccc gcc ctg gag	700		
Ser Leu Pro Ile Leu Arg Pro Ala Gly Thr Gly Pro Pro Ala Leu Glu			
170	175	180	
 cgt gtg gac gcc cag agc cgc cgg gag agc ctg gac atc ttg gcc cct	748		
Arg Val Asp Ala Gln Ser Arg Arg Glu Ser Leu Asp Ile Leu Ala Pro			
185	190	195	
 ggc cgc cgc cgc aag aac atg tcg gag ttc ctg ggg gag gcg agc atc	796		
Gly Arg Arg Arg Lys Asn Met Ser Glu Phe Leu Gly Glu Ala Ser Ile			
200	205	210	215
 ccc ggg cag gag ccc ccc acg ccc tcc agc tgc tct ctg ccc agc ggc	844		
Pro Gly Gln Glu Pro Pro Thr Pro Ser Ser Cys Ser Leu Pro Ser Gly			
220	225	230	
 agc agt ggc agc acc aac act ggc gac agc tgg aag aac cgg gcg gcc	892		

Ser Ser Gly Ser Thr Asn Thr Gly Asp Ser Trp Lys Asn Arg Ala Ala

235

240

245

agt cgc ttc agc ggc ttt ttc agc tcc ggc ccc agc acc acc agc gcc ttt 940

Ser Arg Phe Ser Gly Phe Phe Ser Ser Gly Pro Ser Thr Ser Ala Phe

250

255

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ggc cgg gag gta gac aag atg gag cag ctg gag ggc aag ctg cac acc 988

Gly Arg Glu Val Asp Lys Met Glu Gln Leu Glu Gly Lys Leu His Thr

265

270

275

tac agc ctc ttc ggg ctg ccc agg ctg ccc cgg ggg ctg cgc ttc gac 1036

Tyr Ser Leu Phe Gly Leu Pro Arg Leu Pro Arg Gly Leu Arg Phe Asp

280

285

290

295

cat gac tcc tgg gag gag tac gat gaa gac gag gat gag gac aat 1084

His Asp Ser Trp Glu Glu Tyr Asp Glu Asp Glu Asp Asn

300

305

310

gcc tgc ctg agg ctg gag gac agc tgg cgg gag ctc att gat ggg cat 1132

Ala Cys Leu Arg Leu Glu Asp Ser Trp Arg Glu Leu Ile Asp Gly His

315

320

325

gag aag ctg acc cgg cgg cag tgc cac cag cag gag gcg gtg tgg gag 1180

Glu Lys Leu Thr Arg Arg Gln Cys His Gln Gln Glu Ala Val Trp Glu

330

335

340

ctg ctg cac acg gag gcc tcc tac atc agg aaa ctg cgg gtg atc atc 1228

Leu Leu His Thr Glu Ala Ser Tyr Ile Arg Lys Leu Arg Val Ile Ile

345	350	355	
aac ctg ttc ctg tgc tgc ctc ctg aac ctg caa gag tca ggg ctg ctg 1276			
Asn Leu Phe Leu Cys Cys Leu Leu Asn Leu Gln Glu Ser Gly Leu Leu			
360	365	370	375
tgt gag gtg gag gcg gag cgc ctg ttc agc aac atc ccg gag atc gcg 1324			
Cys Glu Val Glu Ala Glu Arg Leu Phe Ser Asn Ile Pro Glu Ile Ala			
380	385	390	
cag ctg cac cgc agg ctg tgg gct agc gtg atg gcg ccg gtg ctg gag 1372			
Gln Leu His Arg Arg Leu Trp Ala Ser Val Met Ala Pro Val Leu Glu			
395	400	405	
aag gcg cgg cgc acg cga gcg ctg cta cag ccc ggg gac ttc ctc aaa 1420			
Lys Ala Arg Arg Thr Arg Ala Leu Leu Gln Pro Gly Asp Phe Leu Lys			
410	415	420	
ggc ttc aag atg ttc ggc tcg ctc ttc aag ccc tac atc cgc tac tgc 1468			
Gly Phe Lys Met Phe Gly Ser Leu Phe Lys Pro Tyr Ile Arg Tyr Cys			
425	430	435	
atg gag gag ggc tgc atg gag tac atg cgc ggc ctg ctg cgc gac 1516			
Met Glu Glu Glu Gly Cys Met Glu Tyr Met Arg Gly Leu Leu Arg Asp			
440	445	450	455
aac gac ctc ttc cgg gcc tac atc acg tgg gcg gag aag cac cca cag 1564			
Asn Asp Leu Phe Arg Ala Tyr Ile Thr Trp Ala Glu Lys His Pro Gln			
460	465	470	

tgc cag agg ctg aag ctg agc gac atg ctg gcc aaa ccc cac cag cgg	1612		
Cys Gln Arg Leu Lys Leu Ser Asp Met Leu Ala Lys Pro His Gln Arg			
475	480	485	
ctc acc aag tac ccg ctg ctg ctc aag tcg gtg ctg agg aag acc gag	1660		
Leu Thr Lys Tyr Pro Leu Leu Lys Ser Val Leu Arg Lys Thr Glu			
490	495	500	
gag ccg cgc gcc aag gag gcc gtc gtc gcc atg atc ggc tcc gtg gag	1708		
Glu Pro Arg Ala Lys Glu Ala Val Val Ala Met Ile Gly Ser Val Glu			
505	510	515	
cgc ttc atc cac cac gtg aac gcg tgc atg cgg cag cgg cag gag cgg	1756		
Arg Phe Ile His His Val Asn Ala Cys Met Arg Gln Arg Gln Glu Arg			
520	525	530	535
cag cgg ctg gcg gcc gtg gtg agc cgc atc gac gcc tac gag gtg gtg	1804		
Gln Arg Leu Ala Ala Val Val Ser Arg Ile Asp Ala Tyr Glu Val Val			
540	545	550	
gaa agc agc agc gac gaa gtg gac aag ctc ctg aag gaa ttt ctg cac	1852		
Glu Ser Ser Ser Asp Glu Val Asp Lys Leu Leu Lys Glu Phe Leu His			
555	560	565	
ctg gac ttg aca gcg ccc atc cct ggc gcc tcc ccg gag gag acg cgg	1900		
Leu Asp Leu Thr Ala Pro Ile Pro Gly Ala Ser Pro Glu Glu Thr Arg			
570	575	580	

cag ctg ctg ctg gag ggg agc ctg agg atg aag gag ggg aag gac agc			1948
Gln Leu Leu Leu Glu Gly Ser Leu Arg Met Lys Glu Gly Lys Asp Ser			
585	590	595	
aag atg gat gtg tac tgc ttc ctc ttc acg gat ctg ctg ttg gtg acc			1996
Lys Met Asp Val Tyr Cys Phe Leu Phe Thr Asp Leu Leu Leu Val Thr			
600	605	610	615
aaa gca gtg aag aag gca gag agg acc agg gtc atc agg cca ccc ctg			2044
Lys Ala Val Lys Lys Ala Glu Arg Thr Arg Val Ile Arg Pro Pro Leu			
620	625	630	
ctc gtg gac aag att gtg tgc cgg gag cta cgg gac cct ggg tcc ttc			2092
Leu Val Asp Lys Ile Val Cys Arg Glu Leu Arg Asp Pro Gly Ser Phe			
635	640	645	
ctc ctt atc tac ctg aat gag ttt cac agt gct gta ggg gcc tac acg			2140
Leu Leu Ile Tyr Leu Asn Glu Phe His Ser Ala Val Gly Ala Tyr Thr			
650	655	660	
ttc cag gcc agt ggc cag gcc ttg tgc cgt ggc tgg gtg gac acc att			2188
Phe Gln Ala Ser Gly Gln Ala Leu Cys Arg Gly Trp Val Asp Thr Ile			
665	670	675	
tac aat gcc cag aac cag cag ctg caa cag ctg cgt gca cag gag ccc cca			2236
Tyr Asn Ala Gln Asn Gln Leu Gln Gln Leu Arg Ala Gln Glu Pro Pro			
680	685	690	695
ggc agt cag cag ccc ctg cag agc ctg gaa gag gag gag gat gag cag			2284

Gly Ser Gln Gln Pro Leu Gln Ser Leu Glu Glu Glu Asp Glu Gln

700

705

710

gag gag gaa gag gag gag gag gag gag gag gaa ggc gag gac agt 2332

Glu Glu Glu Glu Glu Glu Glu Glu Glu Gly Glu Asp Ser

715

720

725

ggc act tca gct gcc agc tcc cct acc atc atg cgg aaa agc agc ggc 2380

Gly Thr Ser Ala Ala Ser Ser Pro Thr Ile Met Arg Lys Ser Ser Gly

730

735

740

agc ccc gac tct cag cac tgt gcc tca gat ggc tcc acg gag acc ctg 2428

Ser Pro Asp Ser Gln His Cys Ala Ser Asp Gly Ser Thr Glu Thr Leu

745

750

755

gcc atg gtt gtg gta gag cct ggg gac acg ctg tcc tcc ccc gag ttc 2476

Ala Met Val Val Val Glu Pro Gly Asp Thr Leu Ser Ser Pro Glu Phe

760

765

770

775

gac agc ggt cct ttc agc tcc cag tct gat gag acc tct ctc agc acc 2524

Asp Ser Gly Pro Phe Ser Ser Gln Ser Asp Glu Thr Ser Leu Ser Thr

780

785

790

act gcc tca tct gcc acg ccc acc agt gag ctg ctg ccc ctg ggt ccg 2572

Thr Ala Ser Ser Ala Thr Pro Thr Ser Glu Leu Leu Pro Leu Gly Pro

795

800

805

gtg gac ggc cgc tcc tgc tcc atg gac tct gcc tac ggc acc ctc tcc 2620

Val Asp Gly Arg Ser Cys Ser Met Asp Ser Ala Tyr Gly Thr Leu Ser

810	815	820	
cca acc tcc tta caa gac ttt gtg gcc cca ggc cca atg gca gag cta			2668
Pro Thr Ser Leu Gln Asp Phe Val Ala Pro Gly Pro Met Ala Glu Leu			
825	830	835	
gtg cct cgg gcc cca gag tcc cca cga gtt cct tcc cct cca ccc tcg			2716
Val Pro Arg Ala Pro Glu Ser Pro Arg Val Pro Ser Pro Pro Ser			
840	845	850	855
ccc cgt ctc cgc cgc acc cct gtc cag ctg ttg agc tgc ccg ccc			2764
Pro Arg Leu Arg Arg Arg Thr Pro Val Gln Leu Leu Ser Cys Pro Pro			
860	865	870	
cac ctg ctc aag tct aag tcc gag gcc agc ctc ctc cag ctg ctg gca			2812
His Leu Leu Lys Ser Lys Ser Glu Ala Ser Leu Leu Gln Leu Leu Ala			
875	880	885	
ggg gct ggc acc cat ggg aca ccc tct gcc ccc agc cgc agc ctg tca			2860
Gly Ala Gly Thr His Gly Thr Pro Ser Ala Pro Ser Arg Ser Leu Ser			
890	895	900	
gag ctc tgc ctg gct gtt cca gcc cca ggt att agg act cag ggc tcc			2908
Glu Leu Cys Leu Ala Val Pro Ala Pro Gly Ile Arg Thr Gln Gly Ser			
905	910	915	
cct cag gaa gct ggg ccc agc tgg gat tgc cga ggg gcc cct agc cct			2956
Pro Gln Glu Ala Gly Pro Ser Trp Asp Cys Arg Gly Ala Pro Ser Pro			
920	925	930	935

ggc agc ggt cct ggg cta gtc ggc tgc ctg gcc ggg gaa cct gca ggc 3004
 Gly Ser Gly Pro Gly Leu Val Gly Cys Leu Ala Gly Glu Pro Ala Gly

940 945 950

tcc cac agg aag agg tgt gga gac ctg ccc tcg ggg gcc tct ccc agg 3052
 Ser His Arg Lys Arg Cys Gly Asp Leu Pro Ser Gly Ala Ser Pro Arg

955 960 965

gtc cag cct gag ccc cca cca ggg gtc tct gcc cag cac agg aag ctg 3100
 Val Gln Pro Glu Pro Pro Gly Val Ser Ala Gln His Arg Lys Leu

970 975 980

acc ctg gcc cag ctc tac cga atc agg acc acc ctg ctg ctt aac tcc 3148
 Thr Leu Ala Gln Leu Tyr Arg Ile Arg Thr Thr Leu Leu Leu Asn Ser
 985 990 995

acg ctc act gcc tcg gag gtc tgagcagagg gaggccccca agagtgccat 3199
 Thr Leu Thr Ala Ser Glu Val
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tgccctgccc ccacacctccc agtcccacag gccaccctcg gcttgggctg ggttctgtga 3559

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Val Leu Ala Arg Asn Val Ser Thr Arg Ser Cys Pro Pro Arg Thr Ser

20

25

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Pro Ala Val Asp Leu Glu Glu Glu Glu Ser Ser Val Asp Gly

35

40

45

Lys Gly Asp Arg Lys Ser Thr Gly Leu Lys Leu Ser Lys Lys Ala

50

55

60

Arg Arg Arg His Thr Asp Asp Pro Ser Lys Glu Cys Phe Thr Leu Lys

65

70

75

80

Phe Asp Leu Asn Val Asp Ile Glu Thr Glu Ile Val Pro Ala Met Lys

85	90	95
Lys Lys Ser Leu Gly Glu Val Leu Leu Pro Val Phe Glu Arg Lys Gly		
100	105	110
Ile Ala Leu Gly Lys Val Asp Ile Tyr Leu Asp Gln Ser Asn Thr Pro		
115	120	125
Leu Ser Leu Thr Phe Glu Ala Tyr Arg Phe Gly Gly His Tyr Leu Arg		
130	135	140
Val Lys Ala Pro Ala Lys Pro Gly Asp Glu Gly Lys Val Glu Gln Gly		
145	150	155
160		
Met Lys Asp Ser Lys Ser Leu Ser Leu Pro Ile Leu Arg Pro Ala Gly		
165	170	175
Thr Gly Pro Pro Ala Leu Glu Arg Val Asp Ala Gln Ser Arg Arg Glu		
180	185	190
Ser Leu Asp Ile Leu Ala Pro Gly Arg Arg Lys Asn Met Ser Glu		
195	200	205
Phe Leu Gly Glu Ala Ser Ile Pro Gly Gln Glu Pro Pro Thr Pro Ser		
210	215	220
Ser Cys Ser Leu Pro Ser Gly Ser Ser Gly Ser Thr Asn Thr Gly Asp		
225	230	235
240		

Ser Trp Lys Asn Arg Ala Ala Ser Arg Phe Ser Gly Phe Phe Ser Ser

245

250

255

Gly Pro Ser Thr Ser Ala Phe Gly Arg Glu Val Asp Lys Met Glu Gln

260

265

270

Leu Glu Gly Lys Leu His Thr Tyr Ser Leu Phe Gly Leu Pro Arg Leu

275

280

285

Pro Arg Gly Leu Arg Phe Asp His Asp Ser Trp Glu Glu Glu Tyr Asp

290

295

300

Glu Asp Glu Asp Glu Asp Asn Ala Cys Leu Arg Leu Glu Asp Ser Trp

305

310

315

320

Arg Glu Leu Ile Asp Gly His Glu Lys Leu Thr Arg Arg Gln Cys His

325

330

335

Gln Gln Glu Ala Val Trp Glu Leu Leu His Thr Glu Ala Ser Tyr Ile

340

345

350

Arg Lys Leu Arg Val Ile Ile Asn Leu Phe Leu Cys Cys Leu Leu Asn

355

360

365

Leu Gln Glu Ser Gly Leu Leu Cys Glu Val Glu Ala Glu Arg Leu Phe

370

375

380

Ser Asn Ile Pro Glu Ile Ala Gln Leu His Arg Arg Leu Trp Ala Ser

385

390

395

400

Val Met Ala Pro Val Leu Glu Lys Ala Arg Arg Thr Arg Ala Leu Leu

405

410

415

Gln Pro Gly Asp Phe Leu Lys Gly Phe Lys Met Phe Gly Ser Leu Phe

420

425

430

Lys Pro Tyr Ile Arg Tyr Cys Met Glu Glu Gly Cys Met Glu Tyr

435

440

445

Met Arg Gly Leu Leu Arg Asp Asn Asp Leu Phe Arg Ala Tyr Ile Thr

450

455

460

Trp Ala Glu Lys His Pro Gln Cys Gln Arg Leu Lys Leu Ser Asp Met

465

470

475

480

Leu Ala Lys Pro His Gln Arg Leu Thr Lys Tyr Pro Leu Leu Leu Lys

485

490

495

Ser Val Leu Arg Lys Thr Glu Glu Pro Arg Ala Lys Glu Ala Val Val

500

505

510

Ala Met Ile Gly Ser Val Glu Arg Phe Ile His His Val Asn Ala Cys

515

520

525

Met Arg Gln Arg Gln Glu Arg Gln Arg Leu Ala Ala Val Val Ser Arg

530

535

540

Ile Asp Ala Tyr Glu Val Val Glu Ser Ser Asp Glu Val Asp Lys

545

550

555

560

Leu Leu Lys Glu Phe Leu His Leu Asp Leu Thr Ala Pro Ile Pro Gly

565

570

575

Ala Ser Pro Glu Glu Thr Arg Gln Leu Leu Leu Glu Gly Ser Leu Arg

580

585

590

Met Lys Glu Gly Lys Asp Ser Lys Met Asp Val Tyr Cys Phe Leu Phe

595

600

605

Thr Asp Leu Leu Leu Val Thr Lys Ala Val Lys Lys Ala Glu Arg Thr

610

615

620

Arg Val Ile Arg Pro Pro Leu Leu Val Asp Lys Ile Val Cys Arg Glu

625

630

635

640

Leu Arg Asp Pro Gly Ser Phe Leu Leu Ile Tyr Leu Asn Glu Phe His

645

650

655

Ser Ala Val Gly Ala Tyr Thr Phe Gln Ala Ser Gly Gln Ala Leu Cys

660

665

670

Arg Gly Trp Val Asp Thr Ile Tyr Asn Ala Gln Asn Gln Leu Gln Gln

675

680

685

Leu Arg Ala Gln Glu Pro Pro Gly Ser Gln Gln Pro Leu Gln Ser Leu

690

695

700

Glu Glu Glu Glu Asp Glu Gln Glu Glu Glu Glu Glu Glu Glu
 705 710 715 720

Glu Glu Glu Gly Glu Asp Ser Gly Thr Ser Ala Ala Ser Ser Pro Thr
 725 730 735

Ile Met Arg Lys Ser Ser Gly Ser Pro Asp Ser Gln His Cys Ala Ser
 740 745 750

Asp Gly Ser Thr Glu Thr Leu Ala Met Val Val Val Glu Pro Gly Asp
 755 760 765

Thr Leu Ser Ser Pro Glu Phe Asp Ser Gly Pro Phe Ser Ser Gln Ser
 770 775 780

Asp Glu Thr Ser Leu Ser Thr Thr Ala Ser Ser Ala Thr Pro Thr Ser
 785 790 795 800

Glu Leu Leu Pro Leu Gly Pro Val Asp Gly Arg Ser Cys Ser Met Asp
 805 810 815

Ser Ala Tyr Gly Thr Leu Ser Pro Thr Ser Leu Gln Asp Phe Val Ala
 820 825 830

Pro Gly Pro Met Ala Glu Leu Val Pro Arg Ala Pro Glu Ser Pro Arg
 835 840 845

Val Pro Ser Pro Pro Pro Ser Pro Arg Leu Arg Arg Arg Thr Pro Val
 850 855 860

Gln Leu Leu Ser Cys Pro Pro His Leu Leu Lys Ser Lys Ser Glu Ala

865 870 875 880

Ser Leu Leu Gln Leu Leu Ala Gly Ala Gly Thr His Gly Thr Pro Ser

885 890 895

Ala Pro Ser Arg Ser Leu Ser Glu Leu Cys Leu Ala Val Pro Ala Pro

900 905 910

Gly Ile Arg Thr Gln Gly Ser Pro Gln Glu Ala Gly Pro Ser Trp Asp

915 920 925

Cys Arg Gly Ala Pro Ser Pro Gly Ser Gly Pro Gly Leu Val Gly Cys

930 935 940

Leu Ala Gly Glu Pro Ala Gly Ser His Arg Lys Arg Cys Gly Asp Leu

945 950 955 960

Pro Ser Gly Ala Ser Pro Arg Val Gln Pro Glu Pro Pro Pro Gly Val

965 970 975

Ser Ala Gln His Arg Lys Leu Thr Leu Ala Gln Leu Tyr Arg Ile Arg

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cactggcgcg atgcgggccg tcctctcgcc tg atg ggt tgg aag ccc agc gag 113

Met Gly Trp Lys Pro Ser Glu

1 5

gct aga ggc cag tcc caa agt ctc cag gca tca ggg ctg cag ccc agg 161

Ala Arg Gly Gln Ser Gln Ser Leu Gln Ala Ser Gly Leu Gln Pro Arg

10 15 20

agc ctc aag gcg gcc cgg cgg gcg act gga cgg ccg gac agg tcc cga 209

Ser Leu Lys Ala Ala Arg Arg Ala Thr Gly Arg Pro Asp Arg Ser Arg

25 30 35

gca gcc ccg ccc aac atg gac cca gac ccc cag gcg ggc gtg cag gtg 257

Ala Ala Pro Pro Asn Met Asp Pro Asp Pro Gln Ala Gly Val Gln Val

40 45 50 55

ggc atg cgg gtg gtg cgc ggc gac tgg aag tgg ggc cag cag gac 305
 Gly Met Arg Val Val Arg Gly Val Asp Trp Lys Trp Gly Gln Gln Asp
 60 65 70

ggc ggc gag ggc ggc gtg ggc acg gtg gtg gag ctt ggc cgc cac ggc 353
 Gly Gly Glu Gly Gly Val Gly Thr Val Val Glu Leu Gly Arg His Gly
 75 80 85

agc ccc tcg aca ccc gac cgc aca gtg gtc gtg cag tgg gac cag ggc 401
 Ser Pro Ser Thr Pro Asp Arg Thr Val Val Val Gln Trp Asp Gln Gly
 90 95 100

acg cgc acc aac tac cgc gcc ggc tac cag ggc gcg cac gac ctg ctg 449
 Thr Arg Thr Asn Tyr Arg Ala Gly Tyr Gln Gly Ala His Asp Leu Leu
 105 110 115

ctg tac gac aac gcc cag atc ggc gtc cgg cac ccc aac atc atc tgt 497
 Leu Tyr Asp Asn Ala Gln Ile Gly Val Arg His Pro Asn Ile Ile Cys
 120 125 130 135

gac tgc tgc aag aag cac ggg ctg cgg ggg atg cgc tgg aag tgc cgt 545
 Asp Cys Cys Lys Lys His Gly Leu Arg Gly Met Arg Trp Lys Cys Arg
 140 145 150

gtg tgc ctg gac tac gac ctc tgc acg cag tgc tac atg cac aac aag 593
 Val Cys Leu Asp Tyr Asp Leu Cys Thr Gln Cys Tyr Met His Asn Lys
 155 160 165

cat gag ctc gcc cac gcc ttc gac cgc tac gag acc gct cac tcg cgc 641

His Glu Leu Ala His Ala Phe Asp Arg Tyr Glu Thr Ala His Ser Arg

170 175 180

cct gtc aca ctg agt ccc cgc cag ggc ctc ccg agg atc cca cta agg 689

Pro Val Thr Leu Ser Pro Arg Gln Gly Leu Pro Arg Ile Pro Leu Arg

185 190 195

ggc atc ttc cag gga gcg aag gtg gtg cga ggc ccc ttc tgg gag tgg 737

Gly Ile Phe Gln Gly Ala Lys Val Val Arg Gly Pro Phe Trp Glu Trp

200 205 210 215

ggc tca cag gat gga ggg gaa ggg aaa ccg ggc cgt gtg gtg gac atc 785

Gly Ser Gln Asp Gly Gly Glu Gly Lys Pro Gly Arg Val Val Asp Ile

220 225 230

cgt ggc tgg gat gtg gag aca ggc cgg agt gtg gcc agc gtg acg tgg 833

Arg Gly Trp Asp Val Glu Thr Gly Arg Ser Val Ala Ser Val Thr Trp

235 240 245

gct gat ggt acc acc aat gtg tac cgt gtg ggc cac aag ggc aag gtg 881

Ala Asp Gly Thr Thr Asn Val Tyr Arg Val Gly His Lys Gly Lys Val

250 255 260

gac ctc aag tgt gtg ggc gag gca gcg ggc ggc ttc tac tac aag gac 929

Asp Leu Lys Cys Val Gly Glu Ala Ala Gly Gly Phe Tyr Tyr Lys Asp

265 270 275

cac ctc cca agg ctc ggc aag ccg gcg gag ctg cag cgc agg gtg agt 977

His Leu Pro Arg Leu Gly Lys Pro Ala Glu Leu Gln Arg Arg Val Ser

280

285

290

295

gct gac agc cag ccc ttc cag cac ggg gac aag gtc aag tgt ctg ctg 1025

Ala Asp Ser Gln Pro Phe Gln His Gly Asp Lys Val Lys Cys Leu Leu

300

305

310

gac act gat gtc ctg cgg gag atg cag gaa ggc cac ggc ggc tgg aac 1073

Asp Thr Asp Val Leu Arg Glu Met Gln Glu Gly His Gly Gly Trp Asn

315

320

325

ccc agg atg gcg gag ttt atc gga cag acg ggc acc gtg cat cgt atc 1121

Pro Arg Met Ala Glu Phe Ile Gly Gln Thr Gly Thr Val His Arg Ile

330

335

340

acg gac cgc ggg gac gtg cgc gtg cag ttc aac cac gag acg cgc tgg 1169

Thr Asp Arg Gly Asp Val Arg Val Gln Phe Asn His Glu Thr Arg Trp

345

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acc ttc cac ccc ggg gcg ctc acc aag cac cac tcc ttc tgg gtg ggc 1217

Thr Phe His Pro Gly Ala Leu Thr Lys His His Ser Phe Trp Val Gly

360

365

370

375

gac gtg gtc cgg gtc atc ggc gac ctt gac aca gtg aag cgg ctg cag 1265

Asp Val Val Arg Val Ile Gly Asp Leu Asp Thr Val Lys Arg Leu Gln

380

385

390

gct ggg cat ggc gag tgg acg gac gac atg gcc cct gcc ctg ggc cgc 1313

Ala Gly His Gly Glu Trp Thr Asp Asp Met Ala Pro Ala Leu Gly Arg

395

400

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gtc	ggg	aag	gtg	gtg	aaa	gtg	ttt	gga	gac	ggg	aac	ctg	cgt	gta	gca	1361
Val	Gly	Lys	Val	Val	Lys	Val	Phe	Gly	Asp	Gly	Asn	Leu	Arg	Val	Ala	
410			415									420				
gtc	gct	ggt	cag	cgg	tgg	acc	ttc	agc	ccc	tcc	tgc	ctg	gtg	gcc	tac	1409
Val	Ala	Gly	Gln	Arg	Trp	Thr	Phe	Ser	Pro	Ser	Cys	Leu	Val	Ala	Tyr	
425			430									435				
cgg	ccc	gag	gag	gat	gcc	aac	ctg	gac	gtg	gcc	gag	cgc	gcc	cgg	gag	1457
Arg	Pro	Glu	Glu	Asp	Ala	Asn	Leu	Asp	Val	Ala	Glu	Arg	Ala	Arg	Glu	
440			445			450							455			
aac	aaa	agc	tca	ctg	agc	gtg	gcc	ctg	gac	aag	ctt	cgg	gcc	cag	aag	1505
Asn	Lys	Ser	Ser	Leu	Ser	Val	Ala	Leu	Asp	Lys	Leu	Arg	Ala	Gln	Lys	
460			465									470				
agt	gac	cca	gag	cac	ccg	gga	agg	ctg	gtg	gtg	gag	gtg	gcf	ctg	ggt	1553
Ser	Asp	Pro	Glu	His	Pro	Gly	Arg	Leu	Val	Val	Glu	Val	Ala	Leu	Gly	
475			480									485				
aac	gca	gcc	cgg	gct	ctg	gac	ctg	ctg	cgg	agg	cgc	cca	gag	caa	gtg	1601
Asn	Ala	Ala	Arg	Ala	Leu	Asp	Leu	Leu	Arg	Arg	Arg	Pro	Glu	Gln	Val	
490			495									500				
gac	acc	aag	aac	caa	ggc	agg	acc	gct	ctg	caa	gtg	gct	gcc	tac	ctg	1649
Asp	Thr	Lys	Asn	Gln	Gly	Arg	Thr	Ala	Leu	Gln	Val	Ala	Ala	Tyr	Leu	
505			510									515				

ggc cag gtg gag ttg ata cgg ctg ctg cta caa gcc agg gcg ggc gtg			1697
Gly Gln Val Glu Leu Ile Arg Leu Leu Leu Gln Ala Arg Ala Gly Val			
520	525	530	535
gac ctg ccg gac gac gag ggc aac acg gca ctg cac tac gcg gcc ctg			1745
Asp Leu Pro Asp Asp Glu Gly Asn Thr Ala Leu His Tyr Ala Ala Leu			
540	545	550	
ggg aac cag ccc gag gcc acc agg gtg ctc ctg agt gct ggg tgc cgg			1793
Gly Asn Gln Pro Glu Ala Thr Arg Val Leu Leu Ser Ala Gly Cys Arg			
555	560	565	
gcg gac gcc atc aac agc acc cag agc aca gca ctg cac gtg gcc gtg			1841
Ala Asp Ala Ile Asn Ser Thr Gln Ser Thr Ala Leu His Val Ala Val			
570	575	580	
cag agg ggc ttc ctg gag gtg gtg cgg gcc ctg tgt gag cgc ggc tgt			1889
Gln Arg Gly Phe Leu Glu Val Val Arg Ala Leu Cys Glu Arg Gly Cys			
585	590	595	
gac gtc aac ctg ccc gac gcc cac tcg gac acg ccc ctg cac tcc gcc			1937
Asp Val Asn Leu Pro Asp Ala His Ser Asp Thr Pro Leu His Ser Ala			
600	605	610	615
atc tcg gcg ggc act gga gcc agc ggc att gtc gag gtc ctc acg gag			1985
Ile Ser Ala Gly Thr Gly Ala Ser Gly Ile Val Glu Val Leu Thr Glu			
620	625	630	
gtg cca aac atc gat gtt acc gcc acc aac agc cag ggt ttc acc ctg			2033

Val Pro Asn Ile Asp Val Thr Ala Thr Asn Ser Gln Gly Phe Thr Leu

635

640

645

ctg cac cat gcc tcc ctc aag ggt cac gcg cta gct gtg aga aag att 2081

Leu His His Ala Ser Leu Lys Gly His Ala Leu Ala Val Arg Lys Ile

650

655

660

ctg gct cgg gcg cgg cag ctg gtg gac gcc aag aag gag gac ggc ttc 2129

Leu Ala Arg Ala Arg Gln Leu Val Asp Ala Lys Glu Asp Gly Phe

665

670

675

acg gcg ctg cat ctg gct gcc ctc aac aac cac cgc gag gtg gcc cag 2177

Thr Ala Leu His Leu Ala Ala Leu Asn Asn His Arg Glu Val Ala Gln

680

685

690

695

atc ctc atc cgg gag ggc cgc tgt gac gtg aac gtg cgc aac cgg aag 2225

Ile Leu Ile Arg Glu Gly Arg Cys Asp Val Asn Val Arg Asn Arg Lys

700

705

710

ctg cag tcc ccg ctg cat ctc gcc gtg caa cag gcc cac gtg ggg ctg 2273

Leu Gln Ser Pro Leu His Leu Ala Val Gln Gln Ala His Val Gly Leu

715

720

725

gtg ccg cta ctg gtg gac gct ggg tgc agt gtc aac gcc gag gac gag 2321

Val Pro Leu Leu Val Asp Ala Gly Cys Ser Val Asn Ala Glu Asp Glu

730

735

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gag ggg gac aca gcc ctg cac gtg gcg ctg cag cgt cat cag ctg ctg 2369

Glu Gly Asp Thr Ala Leu His Val Ala Leu Gln Arg His Gln Leu Leu

745

750

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ccc ctg gtg gct gat ggg gcc ggg ggg gac cca ggg ccc ttg cag ctg 2417

Pro Leu Val Ala Asp Gly Ala Gly Gly Asp Pro Gly Pro Leu Gln Leu

760

765

770

775

ctg tcc agg cta cag gcc tcg ggc ctc ccc ggc agc gcg gag ctg acg 2465

Leu Ser Arg Leu Gln Ala Ser Gly Leu Pro Gly Ser Ala Glu Leu Thr

780

785

790

gtg ggc gcg gcg gtc gcc tgc ttc ctg gcg ctg gag ggc gcc gac gtg 2513

Val Gly Ala Ala Val Ala Cys Phe Leu Ala Leu Glu Gly Ala Asp Val

795

800

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agc tac acc aac cac cgc ggt cgg agc ccg ctg gac ctg gcc gcc gag 2561

Ser Tyr Thr Asn His Arg Gly Arg Ser Pro Leu Asp Leu Ala Ala Glu

810

815

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ggt cgc gtg ctc aag gcc ctt cag ggc tgc gcc cag cgc ttc cgg gag 2609

Gly Arg Val Leu Lys Ala Leu Gln Gly Cys Ala Gln Arg Phe Arg Glu

825

830

835

cgg cag gcg ggc ggg ggc gcg gcc ccg ggc ccc agg caa acg ctc ggg 2657

Arg Gln Ala Gly Gly Ala Ala Pro Gly Pro Arg Gln Thr Leu Gly

840

845

850

855

acc ccc aac acc gtg acg aac ctg cac gtg ggc gcc gcg ccc ggg ccc 2705

Thr Pro Asn Thr Val Thr Asn Leu His Val Gly Ala Ala Pro Gly Pro

860

865

870

gag gcc gct gag tgc ctg gtg tgc tcc gag ctg gcg ctg ctg gtg ctg	2753		
Glu Ala Ala Glu Cys Leu Val Cys Ser Glu Leu Ala Leu Leu Val Leu			
875	880	885	
ttc tcg ccg tgc cag cac cgc acc gtg tgt gag gag tgc gcg cgc agg	2801		
Phe Ser Pro Cys Gln His Arg Thr Val Cys Glu Glu Cys Ala Arg Arg			
890	895	900	
atg aag aag tgc atc agg tgc cag gtg gtc gtc agc aag aaa ctg cgc	2849		
Met Lys Lys Cys Ile Arg Cys Gln Val Val Val Ser Lys Lys Leu Arg			
905	910	915	
cca gac ggc tct gag gtg gcg agc gcc ccc gcc ccc ggc ccg ccg	2897		
Pro Asp Gly Ser Glu Val Ala Ser Ala Ala Pro Ala Pro Gly Pro Pro			
920	925	930	935
cgc cag ctg gtg gag gag ctg cag agc cgc tac cgg cag atg gag gaa	2945		
Arg Gln Leu Val Glu Glu Leu Gln Ser Arg Tyr Arg Gln Met Glu Glu			
940	945	950	
cgc atc acc tgc ccc atc tgc atc gac agg cac atc cgc ctc gtg ttc	2993		
Arg Ile Thr Cys Pro Ile Cys Ile Asp Arg His Ile Arg Leu Val Phe			
955	960	965	
cag tgc ggc cac ggc gca tgc gcc ccc tgc ggc tcc gcg ctc agc gcc	3041		
Gln Cys Gly His Gly Ala Cys Ala Pro Cys Gly Ser Ala Leu Ser Ala			
970	975	980	

tgc ccc atc tgc cgc cag ccc atc cgc gac cgc atc cag atc ttc gtg 3089
 Cys Pro Ile Cys Arg Gln Pro Ile Arg Asp Arg Ile Gln Ile Phe Val
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aaaagaaaga ttctcgat 3168

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<212> PRT

<213> Homo sapiens

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Gly Arg Pro Asp Arg Ser Arg Ala Ala Pro Pro Asn Met Asp Pro Asp
 35 40 45

Pro Gln Ala Gly Val Gln Val Gly Met Arg Val Val Arg Gly Val Asp
 50 55 60

Trp Lys Trp Gly Gln Gln Asp Gly Gly Glu Gly Gly Val Gly Thr Val
 65 70 75 80

Val Glu Leu Gly Arg His Gly Ser Pro Ser Thr Pro Asp Arg Thr Val
 85 90 95

Val Val Gln Trp Asp Gln Gly Thr Arg Thr Asn Tyr Arg Ala Gly Tyr
 100 105 110

Gln Gly Ala His Asp Leu Leu Leu Tyr Asp Asn Ala Gln Ile Gly Val
 115 120 125

Arg His Pro Asn Ile Ile Cys Asp Cys Cys Lys Lys His Gly Leu Arg
 130 135 140

Gly Met Arg Trp Lys Cys Arg Val Cys Leu Asp Tyr Asp Leu Cys Thr
 145 150 155 160

Gln Cys Tyr Met His Asn Lys His Glu Leu Ala His Ala Phe Asp Arg
 165 170 175

Tyr Glu Thr Ala His Ser Arg Pro Val Thr Leu Ser Pro Arg Gln Gly
 180 185 190

Leu Pro Arg Ile Pro Leu Arg Gly Ile Phe Gln Gly Ala Lys Val Val
 195 200 205

Arg Gly Pro Phe Trp Glu Trp Gly Ser Gln Asp Gly Gly Glu Gly Lys
 210 215 220

Pro Gly Arg Val Val Asp Ile Arg Gly Trp Asp Val Glu Thr Gly Arg

225

230

235

240

Ser Val Ala Ser Val Thr Trp Ala Asp Gly Thr Thr Asn Val Tyr Arg

245

250

255

Val Gly His Lys Gly Lys Val Asp Leu Lys Cys Val Gly Glu Ala Ala

260

265

270

Gly Gly Phe Tyr Tyr Lys Asp His Leu Pro Arg Leu Gly Lys Pro Ala

275

280

285

Glu Leu Gln Arg Arg Val Ser Ala Asp Ser Gln Pro Phe Gln His Gly

290

295

300

Asp Lys Val Lys Cys Leu Leu Asp Thr Asp Val Leu Arg Glu Met Gln

305

310

315

320

Glu Gly His Gly Gly Trp Asn Pro Arg Met Ala Glu Phe Ile Gly Gln

325

330

335

Thr Gly Thr Val His Arg Ile Thr Asp Arg Gly Asp Val Arg Val Gln

340

345

350

Phe Asn His Glu Thr Arg Trp Thr Phe His Pro Gly Ala Leu Thr Lys

355

360

365

His His Ser Phe Trp Val Gly Asp Val Val Arg Val Ile Gly Asp Leu

370

375

380

Asp Thr Val Lys Arg Leu Gln Ala Gly His Gly Glu Trp Thr Asp Asp
 385 390 395 400

Met Ala Pro Ala Leu Gly Arg Val Gly Lys Val Val Lys Val Phe Gly
 405 410 415

Asp Gly Asn Leu Arg Val Ala Val Ala Gly Gln Arg Trp Thr Phe Ser
 420 425 430

Pro Ser Cys Leu Val Ala Tyr Arg Pro Glu Glu Asp Ala Asn Leu Asp
 435 440 445

Val Ala Glu Arg Ala Arg Glu Asn Lys Ser Ser Leu Ser Val Ala Leu
 450 455 460

Asp Lys Leu Arg Ala Gln Lys Ser Asp Pro Glu His Pro Gly Arg Leu
 465 470 475 480

Val Val Glu Val Ala Leu Gly Asn Ala Ala Arg Ala Leu Asp Leu Leu
 485 490 495

Arg Arg Arg Pro Glu Gln Val Asp Thr Lys Asn Gln Gly Arg Thr Ala
 500 505 510

Leu Gln Val Ala Ala Tyr Leu Gly Gln Val Glu Leu Ile Arg Leu Leu
 515 520 525

Leu Gln Ala Arg Ala Gly Val Asp Leu Pro Asp Asp Glu Gly Asn Thr
 530 535 540

Ala Leu His Tyr Ala Ala Leu Gly Asn Gln Pro Glu Ala Thr Arg Val

545 550 555 560

Leu Leu Ser Ala Gly Cys Arg Ala Asp Ala Ile Asn Ser Thr Gln Ser

565 570 575

Thr Ala Leu His Val Ala Val Gln Arg Gly Phe Leu Glu Val Val Arg

580 585 590

Ala Leu Cys Glu Arg Gly Cys Asp Val Asn Leu Pro Asp Ala His Ser

595 600 605

Asp Thr Pro Leu His Ser Ala Ile Ser Ala Gly Thr Gly Ala Ser Gly

610 615 620

Ile Val Glu Val Leu Thr Glu Val Pro Asn Ile Asp Val Thr Ala Thr

625 630 635 640

Asn Ser Gln Gly Phe Thr Leu Leu His His Ala Ser Leu Lys Gly His

645 650 655

Ala Leu Ala Val Arg Lys Ile Leu Ala Arg Ala Arg Gln Leu Val Asp

660 665 670

Ala Lys Lys Glu Asp Gly Phe Thr Ala Leu His Leu Ala Ala Leu Asn

675 680 685

Asn His Arg Glu Val Ala Gln Ile Leu Ile Arg Glu Gly Arg Cys Asp

690	695	700
Val Asn Val Arg Asn Arg Lys Leu Gln Ser Pro Leu His Leu Ala Val		
705	710	715
Gln Gln Ala His Val Gly Leu Val Pro Leu Leu Val Asp Ala Gly Cys		
725	730	735
Ser Val Asn Ala Glu Asp Glu Glu Gly Asp Thr Ala Leu His Val Ala		
740	745	750
Leu Gln Arg His Gln Leu Leu Pro Leu Val Ala Asp Gly Ala Gly Gly		
755	760	765
Asp Pro Gly Pro Leu Gln Leu Leu Ser Arg Leu Gln Ala Ser Gly Leu		
770	775	780
Pro Gly Ser Ala Glu Leu Thr Val Gly Ala Ala Val Ala Cys Phe Leu		
785	790	795
Ala Leu Glu Gly Ala Asp Val Ser Tyr Thr Asn His Arg Gly Arg Ser		
805	810	815
Pro Leu Asp Leu Ala Ala Glu Gly Arg Val Leu Lys Ala Leu Gln Gly		
820	825	830
Cys Ala Gln Arg Phe Arg Glu Arg Gln Ala Gly Gly Ala Ala Pro		
835	840	845

Gly Pro Arg Gln Thr Leu Gly Thr Pro Asn Thr Val Thr Asn Leu His

850 855 860

Val Gly Ala Ala Pro Gly Pro Glu Ala Ala Glu Cys Leu Val Cys Ser

865 870 875 880

Glu Leu Ala Leu Leu Val Leu Phe Ser Pro Cys Gln His Arg Thr Val

885 890 895

Cys Glu Glu Cys Ala Arg Arg Met Lys Lys Cys Ile Arg Cys Gln Val

900 905 910

Val Val Ser Lys Lys Leu Arg Pro Asp Gly Ser Glu Val Ala Ser Ala

915 920 925

Ala Pro Ala Pro Gly Pro Pro Arg Gln Leu Val Glu Glu Leu Gln Ser

930 935 940

Arg Tyr Arg Gln Met Glu Glu Arg Ile Thr Cys Pro Ile Cys Ile Asp

945 950 955 960

Arg His Ile Arg Leu Val Phe Gln Cys Gly His Gly Ala Cys Ala Pro

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Asp Arg Ile Gln Ile Phe Val

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cactggcgcg atgcgggccg tcctctcgcc tg atg ggt tgg aag ccc agc gag 113

Met Gly Trp Lys Pro Ser Glu

1 5

gct aga ggc cag tcc caa agt ctc cag gca tca ggg ctg cag ccc agg 161

Ala Arg Gly Gln Ser Gln Ser Leu Gln Ala Ser Gly Leu Gln Pro Arg

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agc ctc aag gcg gcc cgg cgg gcg act gga cgg ccg gac agg tcc cga 209

Ser Leu Lys Ala Ala Arg Arg Ala Thr Gly Arg Pro Asp Arg Ser Arg

25 30 35

gca gcc ccg ccc aac atg gac cca gac ccc cag gcg ggc gtg cag gtg 257

Ala Ala Pro Pro Asn Met Asp Pro Asp Pro Gln Ala Gly Val Gln Val

40

45

50

55

ggc atg cgg gtg gtg cgc ggc gtc gac tgg aag tgg ggc cag cag gac 305

Gly Met Arg Val Val Arg Gly Val Asp Trp Lys Trp Gly Gln Gln Asp

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ggc ggc gag ggc ggc gtg ggc acg gtg gtg gag ctt ggc cgc cac ggc 353

Gly Gly Glu Gly Gly Val Gly Thr Val Val Glu Leu Gly Arg His Gly

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agc ccc tcg aca ccc gac cgc aca gtg gtc gtg cag tgg gac cag ggc 401

Ser Pro Ser Thr Pro Asp Arg Thr Val Val Val Gln Trp Asp Gln Gly

90

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acg cgc acc aac tac cgc gcc ggc tac cag ggc gcg cac gac ctg ctg 449

Thr Arg Thr Asn Tyr Arg Ala Gly Tyr Gln Gly Ala His Asp Leu Leu

105

110

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ctg tac gac aac gcc cag atc ggc gtc cgg cac ccc aac atc atc tgt 497

Leu Tyr Asp Asn Ala Gln Ile Gly Val Arg His Pro Asn Ile Ile Cys

120

125

130

135

gac tgc tgc aag aag cac ggg ctg cgg ggg atg cgc tgg aag tgc cgt 545

Asp Cys Cys Lys Lys His Gly Leu Arg Gly Met Arg Trp Lys Cys Arg

140

145

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gtg tgc ctg gac tac gac ctc tgc acg cag tgc tac atg cac aac aag 593

Val Cys Leu Asp Tyr Asp Leu Cys Thr Gln Cys Tyr Met His Asn Lys

155

160

165

cat gag ctc gcc cac gcc ttc gac cgc tac gag acc gct cac tcg cgc 641

His Glu Leu Ala His Ala Phe Asp Arg Tyr Glu Thr Ala His Ser Arg

170

175

180

cct gtc aca ctg agt ccc cgc cag ggc ctc ccg agg atc cca cta agg 689

Pro Val Thr Leu Ser Pro Arg Gln Gly Leu Pro Arg Ile Pro Leu Arg

185

190

195

ggc atc ttc cag gga gcg aag gtg gtg cga ggc ccc ttc tgg gag tgg 737

Gly Ile Phe Gln Gly Ala Lys Val Val Arg Gly Pro Phe Trp Glu Trp

200

205

210

215

ggc tca cag gat gga ggg gaa ggg aaa ccg ggc cgt gtg gtg gac atc 785

Gly Ser Gln Asp Gly Gly Glu Gly Lys Pro Gly Arg Val Val Asp Ile

220

225

230

cgt ggc tgg gat gtg gag aca ggc cgg agt gtg gcc agc gtg acg tgg 833

Arg Gly Trp Asp Val Glu Thr Gly Arg Ser Val Ala Ser Val Thr Trp

235

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gct gat ggt acc acc aat gtg tac cgt gtg ggc cac aag ggc aag gtg 881

Ala Asp Gly Thr Thr Asn Val Tyr Arg Val Gly His Lys Gly Lys Val

250

255

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gac ctc aag tgt gtg ggc gag gca gcg ggc ggc ttc tac tac aag gac 929

Asp Leu Lys Cys Val Gly Glu Ala Ala Gly Gly Phe Tyr Tyr Lys Asp

265

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275

cac ctc cca agg ctc ggc aag ccg gcg gag ctg cag cgc agg gtg agt 977
 His Leu Pro Arg Leu Gly Lys Pro Ala Glu Leu Gln Arg Arg Val Ser
 280 285 290 295

gct gac agc cag ccc ttc cag cac ggg gac aag gtc aag tgt ctg ctg 1025
 Ala Asp Ser Gln Pro Phe Gln His Gly Asp Lys Val Lys Cys Leu Leu
 300 305 310

gac act gat gtc ctg cgg gag atg cag gaa ggc cac ggc ggc tgg aac 1073
 Asp Thr Asp Val Leu Arg Glu Met Gln Glu Gly His Gly Gly Trp Asn
 315 320 325

ccc agg atg gcg gag ttt atc gga cag acg ggc acc gtg cat cgt atc 1121
 Pro Arg Met Ala Glu Phe Ile Gly Gln Thr Gly Thr Val His Arg Ile
 330 335 340

acg gac cgc ggg gac gtg cgc gtg cag ttc aac cac gag acg cgc tgg 1169
 Thr Asp Arg Gly Asp Val Arg Val Gln Phe Asn His Glu Thr Arg Trp
 345 350 355

acc ttc cac ccc ggg gcg ctc acc aag cac cac tcc ttc tgg gtg ggc 1217
 Thr Phe His Pro Gly Ala Leu Thr Lys His His Ser Phe Trp Val Gly
 360 365 370 375

gac gtg gtc cgg gtc atc ggc gac ctt gac aca gtg aag cgg ctg cag 1265
 Asp Val Val Arg Val Ile Gly Asp Leu Asp Thr Val Lys Arg Leu Gln
 380 385 390

gct ggg cat ggc gag tgg acg gac gac atg gcc cct gcc ctg ggc cgc 1313

Ala Gly His Gly Glu Trp Thr Asp Asp Met Ala Pro Ala Leu Gly Arg

395 400 405

gtc ggg aag gtg gtg aaa gtg ttt gga gac ggg aac ctg cgt gta gca 1361

Val Gly Lys Val Val Lys Val Phe Gly Asp Gly Asn Leu Arg Val Ala

410 415 420

gtc gct ggt cag cgg tgg acc ttc agc ccc tcc tgc ctg gtg gcc tac 1409

Val Ala Gly Gln Arg Trp Thr Phe Ser Pro Ser Cys Leu Val Ala Tyr

425 430 435

cgg ccc gag gag gat gcc aac ctg gac gtg gcc gag cgc gcc cgg gag 1457

Arg Pro Glu Glu Asp Ala Asn Leu Asp Val Ala Glu Arg Ala Arg Glu

440 445 450 455

aac aaa agc tca ctg agc gtg gcc ctg gac aag ctt cgg gcc cag aag 1505

Asn Lys Ser Ser Leu Ser Val Ala Leu Asp Lys Leu Arg Ala Gln Lys

460 465 470

agt gac cca gag cac ccg gga agg ctg gtg gtg gag gtg gcg ctg ggt 1553

Ser Asp Pro Glu His Pro Gly Arg Leu Val Val Glu Val Ala Leu Gly

475 480 485

aac gca gcc cgg gct ctg gac ctg ctg cgg agg cgc cca gag caa gtg 1601

Asn Ala Ala Arg Ala Leu Asp Leu Leu Arg Arg Arg Pro Glu Gln Val

490 495 500

gac acc aag aac caa ggc agg acc gct ctg caa gtg gct gcc tac ctg 1649

Asp Thr Lys Asn Gln Gly Arg Thr Ala Leu Gln Val Ala Ala Tyr Leu

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515

ggc cag gtg gag ttg ata cgg ctg ctg cta caa gcc agg gcg ggc gtg 1697

Gly Gln Val Glu Leu Ile Arg Leu Leu Leu Gln Ala Arg Ala Gly Val

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gac ctg ccg gac gac gag ggc aac acg gca ctg cac tac gcg gcc ctg 1745

Asp Leu Pro Asp Asp Glu Gly Asn Thr Ala Leu His Tyr Ala Ala Leu

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ggg aac cag ccc gag gcc acc agg gtg ctc ctg agt gct ggg tgc cgg 1793

Gly Asn Gln Pro Glu Ala Thr Arg Val Leu Leu Ser Ala Gly Cys Arg

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gcg gac gcc atc aac agc acc cag agc aca gca ctg cac gtg gcc gtg 1841

Ala Asp Ala Ile Asn Ser Thr Gln Ser Thr Ala Leu His Val Ala Val

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580

cag agg ggc ttc ctg gag gtg gtg cgg gcc ctg tgt gag cgc ggc tgt 1889

Gln Arg Gly Phe Leu Glu Val Val Arg Ala Leu Cys Glu Arg Gly Cys

585

590

595

gac gtc aac ctg ccc gac gcc cac tcg gac acg ccc ctg cac tcc gcc 1937

Asp Val Asn Leu Pro Asp Ala His Ser Asp Thr Pro Leu His Ser Ala

600

605

610

615

atc tcg gcg ggc act gga gcc agc ggc att gtc gag gtc ctc acg gag 1985

Ile Ser Ala Gly Thr Gly Ala Ser Gly Ile Val Glu Val Leu Thr Glu

620

625

630

gtg cca aac atc gat gtt acc gcc acc aac agc cag ggt ttc acc ctg 2033

Val Pro Asn Ile Asp Val Thr Ala Thr Asn Ser Gln Gly Phe Thr Leu

635

640

645

ctg cac cat gcc tcc ctc aag ggt cac gcg cta gct gtg aga aag att 2081

Leu His His Ala Ser Leu Lys Gly His Ala Leu Ala Val Arg Lys Ile

650

655

660

ctg gct cgg gcg cgg cag ctg gtg gac gcc aag aag gag gac ggc ttc 2129

Leu Ala Arg Ala Arg Gln Leu Val Asp Ala Lys Lys Glu Asp Gly Phe

665

670

675

acg gcg ctg cat ctg gct gcc ctc aac aac cac cgc gag gtg gcc cag 2177

Thr Ala Leu His Leu Ala Ala Leu Asn Asn His Arg Glu Val Ala Gln

680

685

690

695

atc ctc atc cgg gag ggc cgc tgt gac gtg aac gtg cgc aac cgg aag 2225

Ile Leu Ile Arg Glu Gly Arg Cys Asp Val Asn Val Arg Asn Arg Lys

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705

710

ctg cag tcc ccg ctg cat ctc gcc gtg caa cag gcc cac gtg ggg ctg 2273

Leu Gln Ser Pro Leu His Leu Ala Val Gln Gln Ala His Val Gly Leu

715

720

725

gtg ccg cta ctg gtg gac gct ggg tgc agt gtc aac gcc gag gac gag 2321

Val Pro Leu Leu Val Asp Ala Gly Cys Ser Val Asn Ala Glu Asp Glu

730

735

740

gag ggg gac aca gcc ctg cac gtg gcg ctg cag cgt cat cag ctg ctg 2369
 Glu Gly Asp Thr Ala Leu His Val Ala Leu Gln Arg His Gln Leu Leu
 745 750 755

ccc ctg gtg gct gat ggg gcc ggg ggg gac cca ggg ccc ttg cag ctg 2417
 Pro Leu Val Ala Asp Gly Ala Gly Gly Asp Pro Gly Pro Leu Gln Leu
 760 765 770 775

ctg tcc agg cta cag gcc tcg ggc ctc ccc ggc agc gcg gag ctg acg 2465
 Leu Ser Arg Leu Gln Ala Ser Gly Leu Pro Gly Ser Ala Glu Leu Thr
 780 785 790

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 Val Gly Ala Ala Val Ala Cys Phe Leu Ala Leu Glu Gly Ala Asp Val
 795 800 805

agc tac acc aac cac cgc ggt cgg agc ccg ctg gac ctg gcc gcc gag 2561
 Ser Tyr Thr Asn His Arg Gly Arg Ser Pro Leu Asp Leu Ala Ala Glu
 810 815 820

ggt cgc gtg ctc aag gcc ctt cag ggc tgc gcc cag cgc ttc cgg gag 2609
 Gly Arg Val Leu Lys Ala Leu Gln Gly Cys Ala Gln Arg Phe Arg Glu
 825 830 835

cgg cag gcg ggc ggg ggc gcg gcc ccg ggc ccc agg caa acg ctc ggg 2657
 Arg Gln Ala Gly Gly Ala Ala Pro Gly Pro Arg Gln Thr Leu Gly
 840 845 850 855

acc ccc aac acc gtg acg aac ctg cac gtg ggc gcc gcg ccg ggg ccc 2705

Thr Pro Asn Thr Val Thr Asn Leu His Val Gly Ala Ala Pro Gly Pro

860

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870

gag gcc gct gag tgc ctg gtg tgc tcc gag ctg gcg ctg ctg gtg ctg 2753

Glu Ala Ala Glu Cys Leu Val Cys Ser Glu Leu Ala Leu Leu Val Leu

875

880

885

tcc tcg ccg tgc cag cac cgc acc gtg tgt gag gag tgc gcg cgc agg 2801

Phe Ser Pro Cys Gln His Arg Thr Val Cys Glu Glu Cys Ala Arg Arg

890

895

900

atg aag aag tgc atc agg tgc cag gtg gtc gtc agc aag aaa ctg cgc 2849

Met Lys Lys Cys Ile Arg Cys Gln Val Val Val Ser Lys Lys Leu Arg

905

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915

cca gac ggc tct gag gtg gcg agc gcc gcc ccc gcc ccc ggc ccg ccg 2897

Pro Asp Gly Ser Glu Val Ala Ser Ala Ala Pro Ala Pro Gly Pro Pro

920

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cgc cag ctg gtg gag gag ctg cag agc cgc tac cgg cag atg gag gaa 2945

Arg Gln Leu Val Glu Glu Leu Gln Ser Arg Tyr Arg Gln Met Glu Glu

940

945

950

cgc atc acc tgc ccc atc tgc atc gac agc cac atc cgc ctc gtg ttc 2993

Arg Ile Thr Cys Pro Ile Cys Ile Asp Ser His Ile Arg Leu Val Phe

955

960

965

cag tgc ggc cac ggc gca tgc gcc ccc tgc ggc tcc gcg ctc agc gcc 3041

Gln Cys Gly His Gly Ala Cys Ala Pro Cys Gly Ser Ala Leu Ser Ala

970

975

980

tgc ccc atc tgc cgc cag ccc atc cgc gac cgc atc cag atc ttc gtg 3089

Cys Pro Ile Cys Arg Gln Pro Ile Arg Asp Arg Ile Gln Ile Phe Val

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30

Gly Arg Pro Asp Arg Ser Arg Ala Ala Pro Pro Asn Met Asp Pro Asp

35

40

45

Pro Gln Ala Gly Val Gln Val Gly Met Arg Val Val Arg Gly Val Asp

50

55

60

Trp Lys Trp Gly Gln Gln Asp Gly Gly Glu Gly Val Gly Thr Val
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Val Glu Leu Gly Arg His Gly Ser Pro Ser Thr Pro Asp Arg Thr Val
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Val Val Gln Trp Asp Gln Gly Thr Arg Thr Asn Tyr Arg Ala Gly Tyr
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Gln Gly Ala His Asp Leu Leu Leu Tyr Asp Asn Ala Gln Ile Gly Val
 115 120 125

Arg His Pro Asn Ile Ile Cys Asp Cys Cys Lys Lys His Gly Leu Arg
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Gly Met Arg Trp Lys Cys Arg Val Cys Leu Asp Tyr Asp Leu Cys Thr
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Gln Cys Tyr Met His Asn Lys His Glu Leu Ala His Ala Phe Asp Arg
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Tyr Glu Thr Ala His Ser Arg Pro Val Thr Leu Ser Pro Arg Gln Gly
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Leu Pro Arg Ile Pro Leu Arg Gly Ile Phe Gln Gly Ala Lys Val Val
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Arg Gly Pro Phe Trp Glu Trp Gly Ser Gln Asp Gly Gly Glu Gly Lys
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Pro Gly Arg Val Val Asp Ile Arg Gly Trp Asp Val Glu Thr Gly Arg
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Ser Val Ala Ser Val Thr Trp Ala Asp Gly Thr Thr Asn Val Tyr Arg
 245 250 255

Val Gly His Lys Gly Lys Val Asp Leu Lys Cys Val Gly Glu Ala Ala
 260 265 270

Gly Gly Phe Tyr Tyr Lys Asp His Leu Pro Arg Leu Gly Lys Pro Ala
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Glu Leu Gln Arg Arg Val Ser Ala Asp Ser Gln Pro Phe Gln His Gly
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Asp Lys Val Lys Cys Leu Leu Asp Thr Asp Val Leu Arg Glu Met Gln
 305 310 315 320

Glu Gly His Gly Gly Trp Asn Pro Arg Met Ala Glu Phe Ile Gly Gln
 325 330 335

Thr Gly Thr Val His Arg Ile Thr Asp Arg Gly Asp Val Arg Val Gln
 340 345 350

Phe Asn His Glu Thr Arg Trp Thr Phe His Pro Gly Ala Leu Thr Lys
 355 360 365

His His Ser Phe Trp Val Gly Asp Val Val Arg Val Ile Gly Asp Leu

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Asp Gly Asn Leu Arg Val Ala Val Ala Gly Gln Arg Trp Thr Phe Ser		
420	425	430
Pro Ser Cys Leu Val Ala Tyr Arg Pro Glu Glu Asp Ala Asn Leu Asp		
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Val Ala Glu Arg Ala Arg Glu Asn Lys Ser Ser Leu Ser Val Ala Leu		
450	455	460
Asp Lys Leu Arg Ala Gln Lys Ser Asp Pro Glu His Pro Gly Arg Leu		
465	470	475
Val Val Glu Val Ala Leu Gly Asn Ala Ala Arg Ala Leu Asp Leu Leu		
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Arg Arg Arg Pro Glu Gln Val Asp Thr Lys Asn Gln Gly Arg Thr Ala		
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Thr Ala Leu His Val Ala Val Gln Arg Gly Phe Leu Glu Val Val Arg

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Ala Leu Cys Glu Arg Gly Cys Asp Val Asn Leu Pro Asp Ala His Ser

595 600 605

Asp Thr Pro Leu His Ser Ala Ile Ser Ala Gly Thr Gly Ala Ser Gly

610 615 620

Ile Val Glu Val Leu Thr Glu Val Pro Asn Ile Asp Val Thr Ala Thr

625 630 635 640

Asn Ser Gln Gly Phe Thr Leu Leu His His Ala Ser Leu Lys Gly His

645 650 655

Ala Leu Ala Val Arg Lys Ile Leu Ala Arg Ala Arg Gln Leu Val Asp

660 665 670

Ala Lys Lys Glu Asp Gly Phe Thr Ala Leu His Leu Ala Ala Leu Asn

675 680 685

Asn His Arg Glu Val Ala Gln Ile Leu Ile Arg Glu Gly Arg Cys Asp

690 695 700

Val Asn Val Arg Asn Arg Lys Leu Gln Ser Pro Leu His Leu Ala Val

705 710 715 720

Gln Gln Ala His Val Gly Leu Val Pro Leu Leu Val Asp Ala Gly Cys

725 730 735

Ser Val Asn Ala Glu Asp Glu Glu Gly Asp Thr Ala Leu His Val Ala

740 745 750

Leu Gln Arg His Gln Leu Leu Pro Leu Val Ala Asp Gly Ala Gly Gly

755 760 765

Asp Pro Gly Pro Leu Gln Leu Leu Ser Arg Leu Gln Ala Ser Gly Leu

770 775 780

Pro Gly Ser Ala Glu Leu Thr Val Gly Ala Ala Val Ala Cys Phe Leu

785 790 795 800

Ala Leu Glu Gly Ala Asp Val Ser Tyr Thr Asn His Arg Gly Arg Ser

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Pro Leu Asp Leu Ala Ala Glu Gly Arg Val Leu Lys Ala Leu Gln Gly

820 825 830

Cys Ala Gln Arg Phe Arg Glu Arg Gln Ala Gly Gly Ala Ala Pro

835

840

845

Gly Pro Arg Gln Thr Leu Gly Thr Pro Asn Thr Val Thr Asn Leu His

850

855

860

Val Gly Ala Ala Pro Gly Pro Glu Ala Ala Glu Cys Leu Val Cys Ser

865

870

875

880

Glu Leu Ala Leu Leu Val Leu Phe Ser Pro Cys Gln His Arg Thr Val

885

890

895

Cys Glu Glu Cys Ala Arg Arg Met Lys Lys Cys Ile Arg Cys Gln Val

900

905

910

Val Val Ser Lys Lys Leu Arg Pro Asp Gly Ser Glu Val Ala Ser Ala

915

920

925

Ala Pro Ala Pro Gly Pro Pro Arg Gln Leu Val Glu Glu Leu Gln Ser

930

935

940

Arg Tyr Arg Gln Met Glu Glu Arg Ile Thr Cys Pro Ile Cys Ile Asp

945

950

955

960

Ser His Ile Arg Leu Val Phe Gln Cys Gly His Gly Ala Cys Ala Pro

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Asp Arg Ile Gln Ile Phe Val

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Met Ile Ala Trp Arg Leu Pro Leu

1 5

tgc gtg ctc ttg gtg gcc tcc gtc gag agc cac ctg ggg gcc ctg ggg 162

Cys Val Leu Leu Val Ala Ser Val Glu Ser His Leu Gly Ala Leu Gly

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ccc aag aac gtc tcg cag aaa gac gcg gag ttt gag cgc acc tac gcg 210

Pro Lys Asn Val Ser Gln Lys Asp Ala Glu Phe Glu Arg Thr Tyr Ala

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gac gac gtc aac agc gag ctg gtc aac atc tac acc ttc aac cac acc 258
 Asp Asp Val Asn Ser Glu Leu Val Asn Ile Tyr Thr Phe Asn His Thr
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gtg acc cgc aac cg^g acc gag ggt gtg cga gtg tct gtg aat gtc ctg 306
 Val Thr Arg Asn Arg Thr Glu Gly Val Arg Val Ser Val Asn Val Leu
 60 65 70

aac aag cag aaa ggg gc^g cct ttg ctg ttc gtg gtc cgc cag aag gag 354
 Asn Lys Gln Lys Gly Ala Pro Leu Leu Phe Val Val Arg Gln Lys Glu
 75 80 85

gct gtt gtg tcc ttc cag gtg ccc cta atc ctt cga gga ctg tat cag 402
 Ala Val Val Ser Phe Gln Val Pro Leu Ile Leu Arg Gly Leu Tyr Gln
 90 95 100

cg^g aag tac ctc tac caa aaa gtg gaa cga act ctg tgt cag ccc ccc 450
 Arg Lys Tyr Leu Tyr Gln Lys Val Glu Arg Thr Leu Cys Gln Pro Pro
 105 110 115 120

acc aag aat gag tct gag atc cag ttt ttc tat gtg gac gtg tct acc 498
 Thr Lys Asn Glu Ser Glu Ile Gln Phe Phe Tyr Val Asp Val Ser Thr
 125 130 135

ctg tca ccc gtc aat acc act tac cag ctc cga gtc aac cgt gtg gac 546
 Leu Ser Pro Val Asn Thr Thr Tyr Gln Leu Arg Val Asn Arg Val Asp
 140 145 150

aat ttt gtg ctc agg act gga gag ctg ttt acc ttt aat acc act gca 594

Asn Phe Val Leu Arg Thr Gly Glu Leu Phe Thr Phe Asn Thr Thr Ala

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gcc cag ccc cag tac ttc aaa tac gag ttt cct gat ggt gtg gac tcg 642

Ala Gln Pro Gln Tyr Phe Lys Tyr Glu Phe Pro Asp Gly Val Asp Ser

170

175

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gta att gtc aag gtg acc tcc aag aag gcc ttc ccc tgc tca gtc atc 690

Val Ile Val Lys Val Thr Ser Lys Lys Ala Phe Pro Cys Ser Val Ile

185

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200

tcc atc cag gat gtc ctg tgc cct gtc tat gat ctg gac aac agt gta 738

Ser Ile Gln Asp Val Leu Cys Pro Val Tyr Asp Leu Asp Asn Ser Val

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gcc ttc att ggc atg tac cag acg atg act aag aag gca gcc atc act 786

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gtg cag cgg aaa gac ttc ccc agc aac agc ttc tat gtg gtg gtg gta 834

Val Gln Arg Lys Asp Phe Pro Ser Asn Ser Phe Tyr Val Val Val Val

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Val Lys Thr Glu Asp Gln Ala Cys Gly Ser Leu Pro Phe Tyr Pro

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ttt gtg gaa gat gag cca gtg gat caa ggg cac cgt cag aaa aca ctg 930

Phe Val Glu Asp Glu Pro Val Asp Gln Gly His Arg Gln Lys Thr Leu

265 270 275 280

tca gtg ctg gtc tct cag gct gtc aca tct gag gcc tat gtt ggt ggg 978
 Ser Val Leu Val Ser Gln Ala Val Thr Ser Glu Ala Tyr Val Gly Gly

285 290 295

atg ctc ttt tgc ctg ggc ata ttc ttg tcc ttc tac ctg ctg act gtg 1026
 Met Leu Phe Cys Leu Gly Ile Phe Leu Ser Phe Tyr Leu Leu Thr Val
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ctg ctg gcc tgt tgg gag aac tgg agg caa agg aag aag acc ttg ctg 1074
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 315 320 325

gtg gcc ata gac cga gcc tgc cca gaa agt ggt cac gct cgg gtc ttg 1122
 Val Ala Ile Asp Arg Ala Cys Pro Glu Ser Gly His Ala Arg Val Leu
 330 335 340

gct gat tca ttt cct ggc agt gcc cct tac gag ggt tac aac tat ggc 1170
 Ala Asp Ser Phe Pro Gly Ser Ala Pro Tyr Glu Gly Tyr Asn Tyr Gly
 345 350 355 360

tcc ttt gaa aat ggt tcc gga tcc act gac ggg ttg gtt gaa agc gca 1218
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 Arg Ser Phe Asp Ala Val Gly Pro Arg Pro Arg Leu Asp Ser Met Ser
 410 415 420

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 Ser Val Glu Glu Asp Asp Tyr Asp Thr Leu Thr Asp Ile Asp Ser Asp
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 Lys Asn Val Ile Arg Thr Lys Gln Tyr Leu Cys Val Ala Asp Leu Ala
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 Arg Lys Asp Lys Arg Val Leu Arg Lys Tyr Gln Ile Tyr Phe Trp
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 475 480 485

gt_g at_c acc tac ca_g ac_g gt_g gt_g aat gtc aca gg_g aac ca_g ga_c at_c 1602
 Val Ile Thr Tyr Gln Thr Val Val Asn Val Thr Gly Asn Gln Asp Ile
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tgc tac tac aac ttc ctc tgt gcc cac ccg ctg ggc aac ctc agc gcc	1650		
Cys Tyr Tyr Asn Phe Leu Cys Ala His Pro Leu Gly Asn Leu Ser Ala			
505	510	515	520
ttc aac aac atc ctc agc aac ttg ggg tac atc ctg ctg ggg ctg ctc	1698		
Phe Asn Asn Ile Leu Ser Asn Leu Gly Tyr Ile Leu Leu Gly Leu Leu			
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ttc ctg ctc atc atc ctg cag cga gag atc aat cat aac cgg gcc ctg	1746		
Phe Leu Leu Ile Ile Leu Gln Arg Glu Ile Asn His Asn Arg Ala Leu			
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ctg cgg aat gac ctc tat gct ctg gag tgt ggg atc ccc aaa cac ttt	1794		
Leu Arg Asn Asp Leu Tyr Ala Leu Glu Cys Gly Ile Pro Lys His Phe			
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ggt ctg ttt tac gcc atg ggc aca gca ctg atg atg gag ggg cta ctt	1842		
Gly Leu Phe Tyr Ala Met Gly Thr Ala Leu Met Met Glu Gly Leu Leu			
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agt gcc tgt tac cac gtc tgc ccc aac tac acc aac ttc cag ttt gat	1890		
Ser Ala Cys Tyr His Val Cys Pro Asn Tyr Thr Asn Phe Gln Phe Asp			
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acc tcc ttc atg tac atg att gct ggc ctc tgc atg ctg aag ctc tac	1938		
Thr Ser Phe Met Tyr Met Ile Ala Gly Leu Cys Met Leu Lys Leu Tyr			
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cag aag cgg cac cca gat atc aac gcc agt gcc tac agt gca tat gcc	1986		

Gln Lys Arg His Pro Asp Ile Asn Ala Ser Ala Tyr Ser Ala Tyr Ala

620

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tgc ttg gcc atc gtc atc ttc ttc tcc gtt ctg ggc gtg gtg ttt ggc 2034

Cys Leu Ala Ile Val Ile Phe Phe Ser Val Leu Gly Val Val Phe Gly

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aaa ggg aac acg gcc ttc tgg att gtc ttc tcc gtc att cac atc atc 2082

Lys Gly Asn Thr Ala Phe Trp Ile Val Phe Ser Val Ile His Ile Ile

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tcc acc ctg ctc ctc agc act cag ctc tat tac atg ggc cgc tgg aag 2130

Ser Thr Leu Leu Leu Ser Thr Gln Leu Tyr Tyr Met Gly Arg Trp Lys

665

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675

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ctg gac ttc ggg atc ttc cgc cgc atc ctc cat gtg ctc tac aca gac 2178

Leu Asp Phe Gly Ile Phe Arg Arg Ile Leu His Val Leu Tyr Thr Asp

685

690

695

tgc atc cgg cag tgc agc ggg ccc ctt tac acg gac cgc atg gtg ctt 2226

Cys Ile Arg Gln Cys Ser Gly Pro Leu Tyr Thr Asp Arg Met Val Leu

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ctg gtc atg ggc aac att atc aac tgg tcg ctg gct gca tac gga ctc 2274

Leu Val Met Gly Asn Ile Ile Asn Trp Ser Leu Ala Ala Tyr Gly Leu

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atc atg cgc ccc aat gac ttt gct tcc tac ttg ctg gca att ggc atc 2322

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Cys Asn Leu Leu Leu Tyr Phe Ala Phe Tyr Ile Ile Met Lys Leu Arg			
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agc ggc gag agg atc aag ctc atc cct ctg ctt tgc atc gtc tgc acc			2418
Ser Gly Glu Arg Ile Lys Leu Ile Pro Leu Leu Cys Ile Val Cys Thr			
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tcc gtg gtc tgg ggc ttc gcg ctc ttc ttc ttc cag gga ctg agc			2466
Ser Val Val Trp Gly Phe Ala Leu Phe Phe Phe Gln Gly Leu Ser			
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Thr Trp Gln Lys Thr Pro Ala Glu Ser Arg Glu His Asn Arg Asp Cys			
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Ile Leu Leu Asp Phe Phe Asp Asp His Asp Ile Trp His Phe Leu Ser			
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Ser Ile Ala Met Phe Gly Ser Phe Leu Val Leu Leu Thr Leu Asp Asp			
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Asp Leu Asp Thr Val Gln Arg Asp Lys Ile Tyr Val Phe			
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Ala Glu Phe Glu Arg Thr Tyr Ala Asp Asp Val Asn Ser Glu Leu Val		
35	40	45
Asn Ile Tyr Thr Phe Asn His Thr Val Thr Arg Asn Arg Thr Glu Gly		
50	55	60
Val Arg Val Ser Val Asn Val Leu Asn Lys Gln Lys Gly Ala Pro Leu		
65	70	75
Leu Phe Val Val Arg Gln Lys Glu Ala Val Val Ser Phe Gln Val Pro		
85	90	95
Leu Ile Leu Arg Gly Leu Tyr Gln Arg Lys Tyr Leu Tyr Gln Lys Val		
100	105	110
Glu Arg Thr Leu Cys Gln Pro Pro Thr Lys Asn Glu Ser Glu Ile Gln		
115	120	125
Phe Phe Tyr Val Asp Val Ser Thr Leu Ser Pro Val Asn Thr Thr Tyr		
130	135	140
Gln Leu Arg Val Asn Arg Val Asp Asn Phe Val Leu Arg Thr Gly Glu		
145	150	155
Leu Phe Thr Phe Asn Thr Thr Ala Ala Gln Pro Gln Tyr Phe Lys Tyr		
165	170	175

Glu Phe Pro Asp Gly Val Asp Ser Val Ile Val Lys Val Thr Ser Lys

180 185 190

Lys Ala Phe Pro Cys Ser Val Ile Ser Ile Gln Asp Val Leu Cys Pro

195 200 205

Val Tyr Asp Leu Asp Asn Ser Val Ala Phe Ile Gly Met Tyr Gln Thr

210 215 220

Met Thr Lys Lys Ala Ala Ile Thr Val Gln Arg Lys Asp Phe Pro Ser

225 230 235 240

Asn Ser Phe Tyr Val Val Val Val Lys Thr Glu Asp Gln Ala Cys

245 250 255

Gly Gly Ser Leu Pro Phe Tyr Pro Phe Val Glu Asp Glu Pro Val Asp

260 265 270

Gln Gly His Arg Gln Lys Thr Leu Ser Val Leu Val Ser Gln Ala Val

275 280 285

Thr Ser Glu Ala Tyr Val Gly Gly Met Leu Phe Cys Leu Gly Ile Phe

290 295 300

Leu Ser Phe Tyr Leu Leu Thr Val Leu Leu Ala Cys Trp Glu Asn Trp

305 310 315 320

Arg Gln Arg Lys Lys Thr Leu Leu Val Ala Ile Asp Arg Ala Cys Pro

325 330 335

Glu Ser Gly His Ala Arg Val Leu Ala Asp Ser Phe Pro Gly Ser Ala

340 345 350

Pro Tyr Glu Gly Tyr Asn Tyr Gly Ser Phe Glu Asn Gly Ser Gly Ser

355 360 365

Thr Asp Gly Leu Val Glu Ser Ala Gly Ser Gly Asp Leu Ser Tyr Ser

370 375 380

Tyr Gln Gly His Asp Gln Phe Lys Arg Arg Leu Pro Ser Gly Gln Met

385 390 395 400

Arg Gln Leu Cys Ile Ala Met Asp Arg Ser Phe Asp Ala Val Gly Pro

405 410 415

Arg Pro Arg Leu Asp Ser Met Ser Ser Val Glu Glu Asp Asp Tyr Asp

420 425 430

Thr Leu Thr Asp Ile Asp Ser Asp Lys Asn Val Ile Arg Thr Lys Gln

435 440 445

Tyr Leu Cys Val Ala Asp Leu Ala Arg Lys Asp Lys Arg Val Leu Arg

450 455 460

Lys Lys Tyr Gln Ile Tyr Phe Trp Asn Ile Ala Thr Ile Ala Val Phe

465 470 475 480

Tyr Ala Leu Pro Val Val Gln Leu Val Ile Thr Tyr Gln Thr Val Val

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Asn Val Thr Gly Asn Gln Asp Ile Cys Tyr Tyr Asn Phe Leu Cys Ala		
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His Pro Leu Gly Asn Leu Ser Ala Phe Asn Asn Ile Leu Ser Asn Leu		
515	520	525
Gly Tyr Ile Leu Leu Gly Leu Leu Phe Leu Leu Ile Ile Leu Gln Arg		
530	535	540
Glu Ile Asn His Asn Arg Ala Leu Leu Arg Asn Asp Leu Tyr Ala Leu		
545	550	555
560		
Glu Cys Gly Ile Pro Lys His Phe Gly Leu Phe Tyr Ala Met Gly Thr		
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Gly Leu Cys Met Leu Lys Leu Tyr Gln Lys Arg His Pro Asp Ile Asn		
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Ser Val Leu Gly Val Val Phe Gly Lys Gly Asn Thr Ala Phe Trp Ile

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Val Phe Ser Val Ile His Ile Ile Ser Thr Leu Leu Leu Ser Thr Gln

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Ser Tyr Leu Leu Ala Ile Gly Ile Cys Asn Leu Leu Leu Tyr Phe Ala

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Phe Tyr Ile Ile Met Lys Leu Arg Ser Gly Glu Arg Ile Lys Leu Ile

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Pro Leu Leu Cys Ile Val Cys Thr Ser Val Val Trp Gly Phe Ala Leu

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Phe Phe Phe Gly Leu Ser Thr Trp Gln Lys Thr Pro Ala Glu

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Ser Arg Glu His Asn Arg Asp Cys Ile Leu Leu Asp Phe Phe Asp Asp

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Leu Leu Val Ala Ser Val Glu Ser His Leu Gly Val Leu Gly Pro Lys

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Gln Asp Val Leu Cys Pro Val Tyr Asp Leu Asp Asn Asn Val Ala Phe

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 Glu Asp Glu Pro Val Asp Gln Gly His Arg Gln Lys Thr Leu Ser Val
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cag aaa acc cct gca gag tc_g agg gag cac aac cg_g gac tgc atc ctc 2513
 Gln Lys Thr Pro Ala Glu Ser Arg Glu His Asn Arg Asp Cys Ile Leu
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Asn Ile Tyr Thr Phe Asn His Thr Val Thr Arg Asn Arg Thr Glu Gly

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Val Arg Val Ser Val Asn Val Leu Asn Lys Gln Lys Gly Ala Pro Leu

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Glu Phe Pro Glu Gly Val Asp Ser Val Ile Val Lys Val Thr Ser Asn

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Val Tyr Asp Leu Asp Asn Asn Val Ala Phe Ile Gly Met Tyr Gln Thr

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Met Thr Lys Lys Ala Ala Ile Thr Val Gln Arg Lys Asp Phe Pro Ser

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Glu Ser Gly His Pro Arg Val Leu Ala Asp Ser Phe Pro Gly Ser Ser

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Pro Tyr Glu Gly Tyr Asn Tyr Gly Ser Phe Glu Asn Val Ser Gly Ser

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Ala Leu Met Met Glu Gly Leu Leu Ser Ala Cys Tyr His Val Cys Pro

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Ala Ser Ala Tyr Ser Ala Tyr Ala Cys Leu Ala Ile Val Ile Phe Phe

625 630 635 640

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Ile Leu His Val Leu Tyr Thr Asp Cys Ile Arg Gln Cys Ser Gly Pro

690 695 700

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705 710 715 720

Trp Ser Leu Ala Ala Tyr Gly Leu Ile Met Arg Pro Asn Asp Phe Ala

725 730 735

Ser Tyr Leu Leu Ala Ile Gly Ile Cys Asn Leu Leu Leu Tyr Phe Ala

740 745 750

Phe Tyr Ile Ile Met Lys Leu Arg Ser Gly Glu Arg Ile Lys Leu Ile

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cagagtaaac ttgaaggcaga tctgtgcatt ctttcctct gcaacaattt gctcgttct 2751

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gggc 2815

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30

Tyr Pro Thr Tyr Tyr Ile Cys Arg Ser Tyr Glu Asp Cys Cys Gly Ser

35

40

45

Arg Cys Cys Val Arg Ala Leu Ser Ile Gln Arg Leu Trp Tyr Phe Trp

50 55 60

Phe Leu Leu Met Met Gly Val Leu Phe Cys Cys Gly Ala Gly Phe Phe

65 70 75 80

Ile Arg Arg Arg Met Tyr Pro Pro Pro Leu Ile Glu Glu Pro Ala Phe

85 90 95

Asn Val Ser Tyr Thr Arg Gln Pro Pro Asn Pro Gly Pro Gly Ala Gln

100 105 110

Gln Pro Gly Pro Pro Tyr Tyr Asp Pro Gly Gly Pro Gly Met Asn

115 120 125

Pro Val Gly Asn Ser Met Ala Met Ala Phe Gln Val Pro Pro Asn Ser

130 135 140

Pro Gln Gly Ser Val Ala Cys Pro Pro Pro Ala Tyr Cys Asn Thr

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Pro Pro Pro Pro Tyr Glu Gln Val Val Lys Ala Lys

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tacccgagtc tcgtttcctc tcagtccatc cacccttcat ggggccagag ccctctctcc 120

agaatctgag cagca atg ccg ttt gct gaa gac aag acc tat aag tat atc 171

Met Pro Phe Ala Glu Asp Lys Thr Tyr Lys Tyr Ile

1

5

10

tgc cgc aat ttc agc aat ttt tgc aat gtg gat gtt gta gag att ctg 219

Cys Arg Asn Phe Ser Asn Phe Cys Asn Val Asp Val Val Glu Ile Leu

15

20

25

cct tac ctg ccc tgc ctc aca gca aga gac cag gat cga ctg cgg gcc 267

Pro Tyr Leu Pro Cys Leu Thr Ala Arg Asp Gln Asp Arg Leu Arg Ala

30

35

40

acc tgc aca ctc tca ggg aac cgg gac acc ctc tgg cat ctc ttc aat 315

Thr Cys Thr Leu Ser Gly Asn Arg Asp Thr Leu Trp His Leu Phe Asn

45

50

55

60

acc ctt cag cgg cgg ccc ggc tgg gtg gag tac ttc att gcg gca ctg 363

Thr Leu Gln Arg Arg Pro Gly Trp Val Glu Tyr Phe Ile Ala Ala Leu

65	70	75
agg ggc tgt gag cta gtt gat ctc gcg gac gaa gtg gcc tct gtc tac 411		
Arg Gly Cys Glu Leu Val Asp Leu Ala Asp Glu Val Ala Ser Val Tyr		
80	85	90
cag agc tac cag cct cgg acc tcg gac cgt ccc cca gac cca ctg gag 459		
Gln Ser Tyr Gln Pro Arg Thr Ser Asp Arg Pro Pro Asp Pro Leu Glu		
95	100	105
cca ccg tca ctt cct gct gag agg cca ggg ccc ccc aca cct gct gcg 507		
Pro Pro Ser Leu Pro Ala Glu Arg Pro Gly Pro Pro Thr Pro Ala Ala		
110	115	120
gcc cac agc atc ccc tac aac agc tgc aga gag aag gag cca agt tac 555		
Ala His Ser Ile Pro Tyr Asn Ser Cys Arg Glu Lys Glu Pro Ser Tyr		
125	130	135
140		
ccc atg cct gtc cag gag acc cag gcg cca gag tcc cca gga gag aat 603		
Pro Met Pro Val Gln Glu Thr Gln Ala Pro Glu Ser Pro Gly Glu Asn		
145	150	155
tca gag caa gcc ctg cag acg ctc agc ccc aga gcc atc cca agg aat 651		
Ser Glu Gln Ala Leu Gln Thr Leu Ser Pro Arg Ala Ile Pro Arg Asn		
160	165	170
cca gat ggt ggc ccc ctg gag tcc tcc tct gac ctg gca gcc ctc agc 699		
Pro Asp Gly Gly Pro Leu Glu Ser Ser Asp Leu Ala Ala Leu Ser		
175	180	185

cct ctg acc tcc agc ggg cat cag gag cag gac aca gaa ctg ggc agt 747
 Pro Leu Thr Ser Ser Gly His Gln Glu Gln Asp Thr Glu Leu Gly Ser
 190 195 200

acc cac aca gca ggt gcg acc tcc agc ctc aca cca tcc cgt ggg cct 795
 Thr His Thr Ala Gly Ala Thr Ser Ser Leu Thr Pro Ser Arg Gly Pro
 205 210 215 220

gtg tct cca tct gtc tcc ttc cag ccc ctg gcc cgt tcc acc ccc agg 843
 Val Ser Pro Ser Val Ser Phe Gln Pro Leu Ala Arg Ser Thr Pro Arg
 225 230 235

gca agc cgc ttg cct gga ccc aca ggg tca gtt gta tct act ggc acc 891
 Ala Ser Arg Leu Pro Gly Pro Thr Gly Ser Val Val Ser Thr Gly Thr
 240 245 250

tcc ttc tcc tcc tca tcc cct ggc ttg gcc tct gca ggg gct gca gag 939
 Ser Phe Ser Ser Ser Pro Gly Leu Ala Ser Ala Gly Ala Ala Glu
 255 260 265

ggt aaa cag ggt gca gag agt gac cag gcc gag cct atc atc tgc tcc 987
 Gly Lys Gln Gly Ala Glu Ser Asp Gln Ala Glu Pro Ile Ile Cys Ser
 270 275 280

agt ggg gca gag gca cct gcc aac tct ctg ccc tcc aaa gtg cct acc 1035
 Ser Gly Ala Glu Ala Pro Ala Asn Ser Leu Pro Ser Lys Val Pro Thr
 285 290 295 300

acc ttg atg cct gtg aac aca gtg gcc ctg aaa gtg cct gcc aac cca 1083
 Thr Leu Met Pro Val Asn Thr Val Ala Leu Lys Val Pro Ala Asn Pro
 305 310 315

gca tct gtc agc aca gtg ccc tcc aag ttg cca act agc tca aag ccc 1131
 Ala Ser Val Ser Thr Val Pro Ser Lys Leu Pro Thr Ser Ser Lys Pro
 320 325 330

cct ggt gca gtg cct tct aat gcg ctc acc aat cca gca cca tcc aaa 1179
 Pro Gly Ala Val Pro Ser Asn Ala Leu Thr Asn Pro Ala Pro Ser Lys
 335 340 345

ttg ccc atc aac tca acc cgt gct ggc atg gtg cca tcc aaa gtg cct 1227
 Leu Pro Ile Asn Ser Thr Arg Ala Gly Met Val Pro Ser Lys Val Pro
 350 355 360

act agc atg gtg ctc acc aag gtg tct gcc agc aca gtc ccc act gac 1275
 Thr Ser Met Val Leu Thr Lys Val Ser Ala Ser Thr Val Pro Thr Asp
 365 370 375 380

ggg agc agc aga aat gag gag acc cca gca gct cca aca ccc gcc ggc 1323
 Gly Ser Ser Arg Asn Glu Glu Thr Pro Ala Ala Pro Thr Pro Ala Gly
 385 390 395

gcc act gga ggc agc tca gcc tgg cta gac agc agc tct gag aat agg 1371
 Ala Thr Gly Gly Ser Ser Ala Trp Leu Asp Ser Ser Ser Glu Asn Arg
 400 405 410

ggc ctt ggg tcg gag ctg agt aag cct ggc gtg ctg gca tcc cag gta 1419

Gly Leu Gly Ser Glu Leu Ser Lys Pro Gly Val Leu Ala Ser Gln Val

415

420

425

gac agc ccg ttc tcg ggc tgc ttc gag gat ctt gcc atc agt gcc agc 1467

Asp Ser Pro Phe Ser Gly Cys Phe Glu Asp Leu Ala Ile Ser Ala Ser

430

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440

acc tcc ttg ggc atg ggg ccc tgc cat ggc cca gag gag aat gag tat 1515

Thr Ser Leu Gly Met Gly Pro Cys His Gly Pro Glu Glu Asn Glu Tyr

445

450

455

460

aag tcc gag ggc acc ttt ggg atc cac gtg gct gag aac ccc agc atc 1563

Lys Ser Glu Gly Thr Phe Gly Ile His Val Ala Glu Asn Pro Ser Ile

465

470

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cag ctc ctg gag ggc aac cct ggg cca cct gcg gac ccg gat ggc ggc 1611

Gln Leu Leu Glu Gly Asn Pro Gly Pro Pro Ala Asp Pro Asp Gly Gly

480

485

490

ccc agg cca caa gcc gac cgg aag ttc cag gag agg gag gtg cca tgc 1659

Pro Arg Pro Gln Ala Asp Arg Lys Phe Gln Glu Arg Glu Val Pro Cys

495

500

505

cac agg ccc tca cct ggg gct ctg tgg ctc cag gtg gct gtg aca ggg 1707

His Arg Pro Ser Pro Gly Ala Leu Trp Leu Gln Val Ala Val Thr Gly

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515

520

gtg ctg gta gtc aca ctc ctg gtg gtg ctg tac cgg cgg cgt ctg cac 1755

Val Leu Val Val Thr Leu Leu Val Val Leu Tyr Arg Arg Arg Leu His

525

530

535

540

tagtgaagcc ctgggcctt cccaccaccc atctgttccg ttccctgcagt atacctggcc 1815

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tctgcaaggg acaagaacta gagccatcg gggctggct cactgtggcc ccaccccaag 3015

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caagtccagc aacctgcacc ctggAACAGC gagtggaccc taccgggct gtctgtatta 3195

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tagtaccca tgctggcag gtcagttac aatttatgca caggtactag tttattgta 3315

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<212> PRT

<213> Homo sapiens

<400> 170

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15

Ser Asn Phe Cys Asn Val Asp Val Val Glu Ile Leu Pro Tyr Leu Pro

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25

30

Cys Leu Thr Ala Arg Asp Gln Asp Arg Leu Arg Ala Thr Cys Thr Leu

35

40

45

Ser Gly Asn Arg Asp Thr Leu Trp His Leu Phe Asn Thr Leu Gln Arg

50

55

60

Arg Pro Gly Trp Val Glu Tyr Phe Ile Ala Ala Leu Arg Gly Cys Glu

65

70

75

80

Leu Val Asp Leu Ala Asp Glu Val Ala Ser Val Tyr Gln Ser Tyr Gln

85

90

95

Pro Arg Thr Ser Asp Arg Pro Pro Asp Pro Leu Glu Pro Pro Ser Leu

100

105

110

Pro Ala Glu Arg Pro Gly Pro Pro Thr Pro Ala Ala Ala His Ser Ile

115

120

125

Pro Tyr Asn Ser Cys Arg Glu Lys Glu Pro Ser Tyr Pro Met Pro Val

130 135 140

Gln Glu Thr Gln Ala Pro Glu Ser Pro Gly Glu Asn Ser Glu Gln Ala

145 150 155 160

Leu Gln Thr Leu Ser Pro Arg Ala Ile Pro Arg Asn Pro Asp Gly Gly

165 170 175

Pro Leu Glu Ser Ser Ser Asp Leu Ala Ala Leu Ser Pro Leu Thr Ser

180 185 190

Ser Gly His Gln Glu Gln Asp Thr Glu Leu Gly Ser Thr His Thr Ala

195 200 205

Gly Ala Thr Ser Ser Leu Thr Pro Ser Arg Gly Pro Val Ser Pro Ser

210 215 220

Val Ser Phe Gln Pro Leu Ala Arg Ser Thr Pro Arg Ala Ser Arg Leu

225 230 235 240

Pro Gly Pro Thr Gly Ser Val Val Ser Thr Gly Thr Ser Phe Ser Ser

245 250 255

Ser Ser Pro Gly Leu Ala Ser Ala Gly Ala Ala Glu Gly Lys Gln Gly

260 265 270

Ala Glu Ser Asp Gln Ala Glu Pro Ile Ile Cys Ser Ser Gly Ala Glu

275 280 285

Ala Pro Ala Asn Ser Leu Pro Ser Lys Val Pro Thr Thr Leu Met Pro

290 295 300

Val Asn Thr Val Ala Leu Lys Val Pro Ala Asn Pro Ala Ser Val Ser

305 310 315 320

Thr Val Pro Ser Lys Leu Pro Thr Ser Ser Lys Pro Pro Gly Ala Val

325 330 335

Pro Ser Asn Ala Leu Thr Asn Pro Ala Pro Ser Lys Leu Pro Ile Asn

340 345 350

Ser Thr Arg Ala Gly Met Val Pro Ser Lys Val Pro Thr Ser Met Val

355 360 365

Leu Thr Lys Val Ser Ala Ser Thr Val Pro Thr Asp Gly Ser Ser Arg

370 375 380

Asn Glu Glu Thr Pro Ala Ala Pro Thr Pro Ala Gly Ala Thr Gly Gly

385 390 395 400

Ser Ser Ala Trp Leu Asp Ser Ser Ser Glu Asn Arg Gly Leu Gly Ser

405 410 415

Glu Leu Ser Lys Pro Gly Val Leu Ala Ser Gln Val Asp Ser Pro Phe

420 425 430

Ser Gly Cys Phe Glu Asp Leu Ala Ile Ser Ala Ser Thr Ser Leu Gly

435

440

445

Met Gly Pro Cys His Gly Pro Glu Glu Asn Glu Tyr Lys Ser Glu Gly

450

455

460

Thr Phe Gly Ile His Val Ala Glu Asn Pro Ser Ile Gln Leu Leu Glu

465

470

475

480

Gly Asn Pro Gly Pro Pro Ala Asp Pro Asp Gly Gly Pro Arg Pro Gln

485

490

495

Ala Asp Arg Lys Phe Gln Glu Arg Glu Val Pro Cys His Arg Pro Ser

500

505

510

Pro Gly Ala Leu Trp Leu Gln Val Ala Val Thr Gly Val Leu Val Val

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gcggggcgcc gacgaggagt gcaggactca ggaagggcga gtgcgcggcga acagagcccg 180

ggaaaggagg cagggcaagg ccgggcttgg gggcaggtgg tccgggcatc cagccttgaa 240

g atg cac aag agg aaa gga ccc ccc gga ccc ccg ggc aga ggc gcc gcg 289

Met His Lys Arg Lys Gly Pro Pro Gly Pro Pro Gly Arg Gly Ala Ala

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5

10

15

gcc gcc cgc cag ctg ggc ctg ctg gtt gac ctc tcc cca gat ggc ctg 337

Ala Ala Arg Gln Leu Gly Leu Leu Val Asp Leu Ser Pro Asp Gly Leu

20

25

30

atg atc cct gag gac ggg gct aac gat gaa gaa ctg gag gct gag ttc 385

Met Ile Pro Glu Asp Gly Ala Asn Asp Glu Glu Leu Glu Ala Glu Phe

35

40

45

ttg gct ttg gtc ggg ggc cag ccc cca gcc ctg gag aag ctc aaa ggc 433

Leu Ala Leu Val Gly Gly Gln Pro Pro Ala Leu Glu Lys Leu Lys Gly

50

55

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aaa ggt ccc ttg ccg atg gag gcc att gag aag atg gcc agc ctg tgc 481

Lys Gly Pro Leu Pro Met Glu Ala Ile Glu Lys Met Ala Ser Leu Cys

65

70

75

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atg aga gac ccg gat gag gat gag gag gag ggg acg gat gag gac gac 529

Met Arg Asp Pro Asp Glu Asp Glu Glu Gly Thr Asp Glu Asp Asp

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90

95

ttg gag gct gat gat gac ctg ctg gcg gag cta aat gag gtc ctt gga 577

Leu Glu Ala Asp Asp Asp Leu Leu Ala Glu Leu Asn Glu Val Leu Gly

100

105

110

gag gag cag aag gct tca gag acc cca cct cct gtg gcc cag ccg aag 625

Glu Glu Gln Lys Ala Ser Glu Thr Pro Pro Pro Val Ala Gln Pro Lys

115

120

125

cct gag gcc cct cat ccg ggg ctg gag acc acc ttg cag gag agg ctg 673

Pro Glu Ala Pro His Pro Gly Leu Glu Thr Thr Leu Gln Glu Arg Leu

130

135

140

gcg ctc tat cag aca gca att gaa agc gcc aga caa gct gga gac agc 721

Ala Leu Tyr Gln Thr Ala Ile Glu Ser Ala Arg Gln Ala Gly Asp Ser

145

150

155

160

gcc aag atg cgg cgc tac gat cgg ggg ctt aaa aca ctg gaa aac ctg 769

Ala Lys Met Arg Arg Tyr Asp Arg Gly Leu Lys Thr Leu Glu Asn Leu

165

170

175

ctc gcc tcc atc cgt aag ggc aat gcc att gac gaa gcg gac atc ccg 817

Leu Ala Ser Ile Arg Lys Gly Asn Ala Ile Asp Glu Ala Asp Ile Pro

180

185

190

ccg cca gtg gcc ata gga aaa ggc ccg gcg tcc acg cct acc tac agc 865

Pro Pro Val Ala Ile Gly Lys Gly Pro Ala Ser Thr Pro Thr Tyr Ser

195

200

205

cct gca ccc acc cag ccg gcc cct aga atc gcg tca gcc cca gag ccc 913

Pro Ala Pro Thr Gln Pro Ala Pro Arg Ile Ala Ser Ala Pro Glu Pro

210

215

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agg gtc acc ctg gag gga cct tct gcc acc gcc cca gcc tca tct cca 961

Arg Val Thr Leu Glu Gly Pro Ser Ala Thr Ala Pro Ala Ser Ser Pro

225

230

235

240

ggc ttg gct aag ccc cag atg ccc cca ggt ccc tgc agc cct ggc cct 1009

Gly Leu Ala Lys Pro Gln Met Pro Pro Gly Pro Cys Ser Pro Gly Pro

245

250

255

ctg gcc cag ttg cag agc cgc cag cgc gac tac aag ctg gct gcc ctc 1057

Leu Ala Gln Leu Gln Ser Arg Gln Arg Asp Tyr Lys Ala Ala Leu

260

265

270

cac gcc aag cag cag gga gat acc act gct gcc gct aga cac ttc cgc 1105

His Ala Lys Gln Gln Gly Asp Thr Thr Ala Ala Arg His Phe Arg

275

280

285

gtg gct aag agc ttt gat gct gtc ttg gag gcc ctg agc cgg ggt gag 1153

Val Ala Lys Ser Phe Asp Ala Val Leu Glu Ala Leu Ser Arg Gly Glu

290

295

300

ccc	gtg	gac	ctc	tcc	tgc	ctg	ccc	cct	cca	ccc	gac	cag	ctg	ccc	cca	1201	
Pro	Val	Asp	Leu	Ser	Cys	Leu	Pro	Pro	Pro	Pro	Asp	Gln	Leu	Pro	Pro		
305															320		
gac	cca	ccg	tca	cca	ccg	tcg	cag	cct	ccg	acc	ccc	gct	acg	gcg	ccc	1249	
Asp	Pro	Pro	Ser	Pro	Pro	Ser	Gln	Pro	Pro	Thr	Pro	Thr	Ala	Thr	Ala	Pro	
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															330	335	
tcc	aca	aca	gag	gtg	ccc	cca	ccc	ccg	agg	acc	ctg	ctg	gag	gcg	ctg	1297	
Ser	Thr	Thr	Glu	Val	Pro	Pro	Pro	Pro	Arg	Thr	Leu	Leu	Glu	Ala	Leu		
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															345	350	
gag	cag	cg	atg	gag	cg	tg	ac	g	cc	g	ca	g	cc	a	ag	1345	
Glu	Gln	Arg	Met	Glu	Arg	Tyr	Gln	Val	Ala	Ala	Ala	Gln	Ala	Lys	Ser		
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aag	ggg	gac	cag	cg	aaa	gct	cga	atg	cac	gag	cg	atc	gtc	a	aa	1393	
Lys	Gly	Asp	Gln	Arg	Lys	Ala	Arg	Met	His	Glu	Arg	Ile	Val	Lys	Gln		
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															375	380	
tac	caa	gat	gcc	atc	cga	gcc	cac	a	ag	gct	gg	cg	gcc	gt	gt	1441	
Tyr	Gln	Asp	Ala	Ile	Arg	Ala	His	Lys	Ala	Gly	Arg	Ala	Val	Asp	Val		
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															390	395	400
gct	gaa	ttg	ccc	gt	ccc	cc	gg	tt	cc	cc	a	tc	c	gg	ct	1489	
Ala	Glu	Leu	Pro	Val	Pro	Pro	Gly	Phe	Pro	Pro	Ile	Gln	Gly	Leu	Glu		
															405		
															410	415	
gcc	acc	a	ag	ccc	acc	c	ag	c	ag	gt	gt	gg	gt	g	act	gg	1537

Ala Thr Lys Pro Thr Gln Gln Ser Leu Val Gly Val Leu Glu Thr Ala

420

425

430

atg aag ctg gcc aac cag gat gaa ggc cca gag gat gaa gag gat gag 1585

Met Lys Leu Ala Asn Gln Asp Glu Gly Pro Glu Asp Glu Glu Asp Glu

435

440

445

gtg cct aag aag cag aac agc cct gtg gcc ccc aca gcc cag ccc aaa 1633

Val Pro Lys Lys Gln Asn Ser Pro Val Ala Pro Thr Ala Gln Pro Lys

450

455

460

gcc cca ccc tca aga act ccc cag tcg gga tca gcc cca aca gcc aaa 1681

Ala Pro Pro Ser Arg Thr Pro Gln Ser Gly Ser Ala Pro Thr Ala Lys

465

470

475

480

gcg ccc ccc aaa gcc aca tcc acc aga gcc cag cag cag ctg gcc ttc 1729

Ala Pro Pro Lys Ala Thr Ser Thr Arg Ala Gln Gln Gln Leu Ala Phe

485

490

495

cta gag ggc cgc aag aag cag ctc ctg cag gcc gca ctg cga gcc aag 1777

Leu Glu Gly Arg Lys Lys Gln Leu Leu Gln Ala Ala Leu Arg Ala Lys

500

505

510

cag aaa aac gac gtg gag ggt gcc aag atg cac ctg cgc caa gcc aag 1825

Gln Lys Asn Asp Val Glu Gly Ala Lys Met His Leu Arg Gln Ala Lys

515

520

525

gga ctg gag cct atg ctg gag gcc tcg cgc aat ggg ctg cct gtg gac 1873

Gly Leu Glu Pro Met Leu Glu Ala Ser Arg Asn Gly Leu Pro Val Asp

530 535 540

atc acc aag gtg ccg cct gcc cct gtc aac aag gac gac ttt gcc ctg 1921

Ile Thr Lys Val Pro Pro Ala Pro Val Asn Lys Asp Asp Phe Ala Leu

545 550 555 560

gtc cag cgg cct ggc ccg ggt ctg tct cag gag gcc gcc cgg cgc tat 1969

Val Gln Arg Pro Gly Pro Gly Leu Ser Gln Glu Ala Ala Arg Arg Tyr

565 570 575

ggt gaa ctc acc aag ctc ata cgg cag cag cac gag atg tgc ctg aac 2017

Gly Glu Leu Thr Lys Leu Ile Arg Gln Gln His Glu Met Cys Leu Asn

580 585 590

cac tca aac caa ttc acc cag ctg ggc aac atc act gaa acc acc aag 2065

His Ser Asn Gln Phe Thr Gln Leu Gly Asn Ile Thr Glu Thr Thr Lys

595 600 605

ttt gaa aag ttg gcg gag gac tgt aag cgg agc atg gac att ctg aag 2113

Phe Glu Lys Leu Ala Glu Asp Cys Lys Arg Ser Met Asp Ile Leu Lys

610 615 620

caa gcc ttc gtc cgg ggt ctc ccc acg ccc acc gcc cgc ttt gag caa 2161

Gln Ala Phe Val Arg Gly Leu Pro Thr Pro Thr Ala Arg Phe Glu Gln

625 630 635 640

agg acc ttc agc gtc atc aag atc ttc cct gac ctc agc agc aac gac 2209

Arg Thr Phe Ser Val Ile Lys Ile Phe Pro Asp Leu Ser Ser Asn Asp

645 650 655

atg ctc ctc ttc atc gtg aag ggc atc aac ttg ccc aca ccc cca gga	2257		
Met Leu Leu Phe Ile Val Lys Gly Ile Asn Leu Pro Thr Pro Pro Gly			
660	665	670	
ctg tcc cct ggc gat ctg gat gtc ttt gtt cgg ttt gac ttc ccc tat	2305		
Leu Ser Pro Gly Asp Leu Asp Val Phe Val Arg Phe Asp Phe Pro Tyr			
675	680	685	
ccc aac gtg gaa gaa gct cag aaa gac aag acc agt gtg atc aag aac	2353		
Pro Asn Val Glu Glu Ala Gln Lys Asp Lys Thr Ser Val Ile Lys Asn			
690	695	700	
aca gac tcc cct gag ttc aag gag cag ttc aaa ctc tgc atc aac cgc	2401		
Thr Asp Ser Pro Glu Phe Lys Glu Gln Phe Lys Leu Cys Ile Asn Arg			
705	710	715	720
agc cac cgt ggc ttc cga agg gcc atc cag acc aag ggc atc aag ttc	2449		
Ser His Arg Gly Phe Arg Arg Ala Ile Gln Thr Lys Gly Ile Lys Phe			
725	730	735	
gaa gtg gtt cac aag ggg ggg ctg ttc aag act gac cgg gtg ctg ggg	2497		
Glu Val Val His Lys Gly Gly Leu Phe Lys Thr Asp Arg Val Leu Gly			
740	745	750	
aca gcc cag ctg aag ctg gat gca ctg gag ata gca tgt gag gtc cgg	2545		
Thr Ala Gln Leu Lys Leu Asp Ala Leu Glu Ile Ala Cys Glu Val Arg			
755	760	765	

gag atc ctt gag gtc ctg gat ggt cgc cgg ccc aca ggg ggg cga ctg	2593		
Glu Ile Leu Glu Val Leu Asp Gly Arg Arg Pro Thr Gly Gly Arg Leu			
770	775	780	
gag gta atg gtc cggtt att cgg gag cca ctg aca gcc cag cag ttg gag	2641		
Glu Val Met Val Arg Ile Arg Glu Pro Leu Thr Ala Gln Gln Leu Glu			
785	790	795	800
acg acg aca gag agg tgg ctg gtc att gac cct gtg ccg gca gct gtg	2689		
Thr Thr Thr Glu Arg Trp Leu Val Ile Asp Pro Val Pro Ala Ala Val			
805	810	815	
ccc aca cag gtt gct ggg ccc aaa ggg aag gcc cct cct gtg cct gcc	2737		
Pro Thr Gln Val Ala Gly Pro Lys Gly Lys Ala Pro Pro Val Pro Ala			
820	825	830	
cct gca agg gag tca ggg aac aga tca gcc cgg ccc ctg cat agc ctc	2785		
Pro Ala Arg Glu Ser Gly Asn Arg Ser Ala Arg Pro Leu His Ser Leu			
835	840	845	
agt gtg ctg gcg ttt gac caa gag cgt ctg gag cgg aag atc ctg gcc	2833		
Ser Val Leu Ala Phe Asp Gln Glu Arg Leu Glu Arg Lys Ile Leu Ala			
850	855	860	
ctc agg cag gcg cgg cgg ccg gtg ccc cca gaa gtg gcc cag cag tac	2881		
Leu Arg Gln Ala Arg Arg Pro Val Pro Pro Glu Val Ala Gln Gln Tyr			
865	870	875	880
cag gac atc atg caa cgc agc cag tgg cag agg gca cag ctg gag cag	2929		

Gln Asp Ile Met Gln Arg Ser Gln Trp Gln Arg Ala Gln Leu Glu Gln

885

890

895

ggg ggt gtg ggc atc cga cgg gaa tac aca gcc cag ctg gag cgg cag 2977

Gly Gly Val Gly Ile Arg Arg Glu Tyr Thr Ala Gln Leu Glu Arg Gln

900

905

910

ctg cag ttc tac acg gag gct gcc cgg cgc ctg ggc aac gat ggc agc 3025

Leu Gln Phe Tyr Thr Glu Ala Ala Arg Arg Leu Gly Asn Asp Gly Ser

915

920

925

agg gat gct gca aag gag gcg ctc tat agg cgg aat ctg gta ggg agt 3073

Arg Asp Ala Ala Lys Glu Ala Leu Tyr Arg Arg Asn Leu Val Gly Ser

930

935

940

gag ctg cag cgg ctc cgc agg tgaggagccc atggggcggg cagccccag 3124

Glu Leu Gln Arg Leu Arg Arg

945

950

aaagcgggca gcaggcccg ataccggaa gagccgacac agccacgaac cagacaagca 3184

gacaatcagc ggacaatcgg ttctggactc acccctcatc cggccccca gccccgccag 3244

agcctccgtg gctgcgggtg ttggAACCA tgcctgccag ccagtatgtc cccctcaccc 3304

aggcctggct gggccctgga gagtcctgtt tgcacagccc aggggtgtcc ggcctctggc 3364

ccgccccgga gcagggaggg cggctggggc caagccccga gggccctgc aagcacttta 3424

cttcctgttc ctccccagcc ttaaccccaa agccctcctg cacccaaag aagccactga 3484

ggctggccga gccacactgt ctccccaggg gcgtcgacct ggcccagctg ggtccccagg 3544

ccagcacatg gaataaaata gccagggcca cactc 3579

<210> 172

<211> 951

<212> PRT

<213> Homo sapiens

<400> 172

Met His Lys Arg Lys Gly Pro Pro Gly Pro Pro Gly Arg Gly Ala Ala

1

5

10

15

Ala Ala Arg Gln Leu Gly Leu Leu Val Asp Leu Ser Pro Asp Gly Leu

20

25

30

Met Ile Pro Glu Asp Gly Ala Asn Asp Glu Glu Leu Glu Ala Glu Phe

35

40

45

Leu Ala Leu Val Gly Gly Gln Pro Pro Ala Leu Glu Lys Leu Lys Gly

50

55

60

Lys Gly Pro Leu Pro Met Glu Ala Ile Glu Lys Met Ala Ser Leu Cys

65

70

75

80

Met Arg Asp Pro Asp Glu Asp Glu Glu Glu Gly Thr Asp Glu Asp Asp

85	90	95
Leu Glu Ala Asp Asp Asp Leu Leu Ala Glu Leu Asn Glu Val Leu Gly		
100	105	110
Glu Glu Gln Lys Ala Ser Glu Thr Pro Pro Pro Val Ala Gln Pro Lys		
115	120	125
Pro Glu Ala Pro His Pro Gly Leu Glu Thr Thr Leu Gln Glu Arg Leu		
130	135	140
Ala Leu Tyr Gln Thr Ala Ile Glu Ser Ala Arg Gln Ala Gly Asp Ser		
145	150	155
160		
Ala Lys Met Arg Arg Tyr Asp Arg Gly Leu Lys Thr Leu Glu Asn Leu		
165	170	175
Leu Ala Ser Ile Arg Lys Gly Asn Ala Ile Asp Glu Ala Asp Ile Pro		
180	185	190
Pro Pro Val Ala Ile Gly Lys Gly Pro Ala Ser Thr Pro Thr Tyr Ser		
195	200	205
Pro Ala Pro Thr Gln Pro Ala Pro Arg Ile Ala Ser Ala Pro Glu Pro		
210	215	220
Arg Val Thr Leu Glu Gly Pro Ser Ala Thr Ala Pro Ala Ser Ser Pro		
225	230	235
240		

Gly Leu Ala Lys Pro Gln Met Pro Pro Gly Pro Cys Ser Pro Gly Pro

245 250 255

Leu Ala Gln Leu Gln Ser Arg Gln Arg Asp Tyr Lys Leu Ala Ala Leu

260 265 270

His Ala Lys Gln Gln Gly Asp Thr Thr Ala Ala Ala Arg His Phe Arg

275 280 285

Val Ala Lys Ser Phe Asp Ala Val Leu Glu Ala Leu Ser Arg Gly Glu

290 295 300

Pro Val Asp Leu Ser Cys Leu Pro Pro Pro Pro Asp Gln Leu Pro Pro

305 310 315 320

Asp Pro Pro Ser Pro Pro Ser Gln Pro Pro Thr Pro Ala Thr Ala Pro

325 330 335

Ser Thr Thr Glu Val Pro Pro Pro Arg Thr Leu Leu Glu Ala Leu

340 345 350

Glu Gln Arg Met Glu Arg Tyr Gln Val Ala Ala Ala Gln Ala Lys Ser

355 360 365

Lys Gly Asp Gln Arg Lys Ala Arg Met His Glu Arg Ile Val Lys Gln

370 375 380

Tyr Gln Asp Ala Ile Arg Ala His Lys Ala Gly Arg Ala Val Asp Val

385 390 395 400

Ala Glu Leu Pro Val Pro Pro Gly Phe Pro Pro Ile Gln Gly Leu Glu

405 410 415

Ala Thr Lys Pro Thr Gln Gln Ser Leu Val Gly Val Leu Glu Thr Ala

420 425 430

Met Lys Leu Ala Asn Gln Asp Glu Gly Pro Glu Asp Glu Asp Glu

435 440 445

Val Pro Lys Lys Gln Asn Ser Pro Val Ala Pro Thr Ala Gln Pro Lys

450 455 460

Ala Pro Pro Ser Arg Thr Pro Gln Ser Gly Ser Ala Pro Thr Ala Lys

465 470 475 480

Ala Pro Pro Lys Ala Thr Ser Thr Arg Ala Gln Gln Leu Ala Phe

485 490 495

Leu Glu Gly Arg Lys Lys Gln Leu Leu Gln Ala Ala Leu Arg Ala Lys

500 505 510

Gln Lys Asn Asp Val Glu Gly Ala Lys Met His Leu Arg Gln Ala Lys

515 520 525

Gly Leu Glu Pro Met Leu Glu Ala Ser Arg Asn Gly Leu Pro Val Asp

530 535 540

Ile Thr Lys Val Pro Pro Ala Pro Val Asn Lys Asp Asp Phe Ala Leu

545	550	555	560
-----	-----	-----	-----

Val Gln Arg Pro Gly Pro Gly Leu Ser Gln Glu Ala Ala Arg Arg Tyr

565	570	575
-----	-----	-----

Gly Glu Leu Thr Lys Leu Ile Arg Gln Gln His Glu Met Cys Leu Asn

580	585	590
-----	-----	-----

His Ser Asn Gln Phe Thr Gln Leu Gly Asn Ile Thr Glu Thr Thr Lys

595	600	605
-----	-----	-----

Phe Glu Lys Leu Ala Glu Asp Cys Lys Arg Ser Met Asp Ile Leu Lys

610	615	620
-----	-----	-----

Gln Ala Phe Val Arg Gly Leu Pro Thr Pro Thr Ala Arg Phe Glu Gln

625	630	635	640
-----	-----	-----	-----

Arg Thr Phe Ser Val Ile Lys Ile Phe Pro Asp Leu Ser Ser Asn Asp

645	650	655
-----	-----	-----

Met Leu Leu Phe Ile Val Lys Gly Ile Asn Leu Pro Thr Pro Pro Gly

660	665	670
-----	-----	-----

Leu Ser Pro Gly Asp Leu Asp Val Phe Val Arg Phe Asp Phe Pro Tyr

675	680	685
-----	-----	-----

Pro Asn Val Glu Glu Ala Gln Lys Asp Lys Thr Ser Val Ile Lys Asn

690	695	700
-----	-----	-----

Thr Asp Ser Pro Glu Phe Lys Glu Gln Phe Lys Leu Cys Ile Asn Arg
705 710 715 720

Ser His Arg Gly Phe Arg Arg Ala Ile Gln Thr Lys Gly Ile Lys Phe
725 730 735

Glu Val Val His Lys Gly Gly Leu Phe Lys Thr Asp Arg Val Leu Gly
740 745 750

Thr Ala Gln Leu Lys Leu Asp Ala Leu Glu Ile Ala Cys Glu Val Arg
755 760 765

Glu Ile Leu Glu Val Leu Asp Gly Arg Arg Pro Thr Gly Gly Arg Leu
770 775 780

Glu Val Met Val Arg Ile Arg Glu Pro Leu Thr Ala Gln Gln Leu Glu
785 790 795 800

Thr Thr Thr Glu Arg Trp Leu Val Ile Asp Pro Val Pro Ala Ala Val
805 810 815

Pro Thr Gln Val Ala Gly Pro Lys Gly Lys Ala Pro Pro Val Pro Ala
820 825 830

Pro Ala Arg Glu Ser Gly Asn Arg Ser Ala Arg Pro Leu His Ser Leu
835 840 845

Ser Val Leu Ala Phe Asp Gln Glu Arg Leu Glu Arg Lys Ile Leu Ala
850 855 860

Leu Arg Gln Ala Arg Arg Pro Val Pro Pro Glu Val Ala Gln Gln Tyr

865 870 875 880

Gln Asp Ile Met Gln Arg Ser Gln Trp Gln Arg Ala Gln Leu Glu Gln

885 890 895

Gly Gly Val Gly Ile Arg Arg Glu Tyr Thr Ala Gln Leu Glu Arg Gln

900 905 910

Leu Gln Phe Tyr Thr Glu Ala Ala Arg Arg Leu Gly Asn Asp Gly Ser

915 920 925

Arg Asp Ala Ala Lys Glu Ala Leu Tyr Arg Arg Asn Leu Val Gly Ser

930 935 940

Glu Leu Gln Arg Leu Arg Arg

945 950

<210> 173

<211> 2796

<212> DNA

<213> Homo sapiens

<220>

<221> CDS

<222> (574)..(1683)

<400> 173

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tagtccttaa caaaggaaaa gcgataaatg taaataagct cacatttca gaatgagcgg 180

tttgcagtaa ggagctgcgg cagcccagag tctgtcttt ttggctggg ctaaccttc 240

cctgtttttt gtttttgtt ttgtttgtt tttgtttttt atggataaaa atatgcgctt 300

ccgaagtgcg agttgccagt ttacacgtt attagctaac tatctacagg catgagcaca 360

ttctctcatc tagcacactc tttcttggc actcaattga ggaactctct gatgtctgc 420

ctccagaaaa ttcattgatt atccaagtct cagataaatc tggtgccaga gtttggtttg 480

aactaactaa tgaagaaagc attctctact ggtcctcagt ctcaagagtg gtgaaccct 540

gcacctagca ggctctctgg gaaaaaaaaa tcc atg ggt gac aga aga ttt att 594

Met Gly Asp Arg Arg Phe Ile

1

5

gac ttc caa ttc caa gat tta aat tca agt ctc aga ccc agg ttg gga 642

Asp Phe Gln Phe Gln Asp Leu Asn Ser Ser Leu Arg Pro Arg Leu Gly

10

15

20

aat gca act gcc aat aat act tgc att gtt gat gat tcc ttc aag tat 690

Asn Ala Thr Ala Asn Asn Thr Cys Ile Val Asp Asp Ser Phe Lys Tyr

25	30	35
----	----	----

aat ttg aat ggt gct gtc tat agt gtt gta ttc atc ctg ggt cta ata 738

Asn Leu Asn Gly Ala Val Tyr Ser Val Val Phe Ile Leu Gly Leu Ile

40	45	50	55
----	----	----	----

acc aac agt gcc tcc ctg ttt gtc ttc tgc ttc cgc atg aaa atg aga 786

Thr Asn Ser Ala Ser Leu Phe Val Phe Cys Phe Arg Met Lys Met Arg

60	65	70
----	----	----

agt gag acg gct act ttc atc acc aac ctg gcc ctc tct gat ttg ctt 834

Ser Glu Thr Ala Thr Phe Ile Thr Asn Leu Ala Leu Ser Asp Leu Leu

75	80	85
----	----	----

ttt gtt tgt acc cta cct ttc aaa ata ttt tac aac ttt aat cgc cac 882

Phe Val Cys Thr Leu Pro Phe Lys Ile Phe Tyr Asn Phe Asn Arg His

90	95	100
----	----	-----

tgg cct ttt ggt gac acc ctc tgt aag atc tca ggg act gcg ttc ctc 930

Trp Pro Phe Gly Asp Thr Leu Cys Lys Ile Ser Gly Thr Ala Phe Leu

105	110	115
-----	-----	-----

acc aac atc tat ggg agc atg ctc ttc ctc acc tgc atc agt gtg gat 978

Thr Asn Ile Tyr Gly Ser Met Leu Phe Leu Thr Cys Ile Ser Val Asp

120	125	130	135
-----	-----	-----	-----

cgt ttc cta gcc att gtc tat ccc ttc cga tcg cgt acc atc agg acc 1026

Arg Phe Leu Ala Ile Val Tyr Pro Phe Arg Ser Arg Thr Ile Arg Thr

140	145	150	
agg agg aat tcc gcc att gtg tgc gct gga gtc tgg atc cta gtc ctc			1074
Arg Arg Asn Ser Ala Ile Val Cys Ala Gly Val Trp Ile Leu Val Leu			
155	160	165	
agt ggt ggt att tca gct tct ttg ttc tcc acc act aat gtc aac aat			1122
Ser Gly Gly Ile Ser Ala Ser Leu Phe Ser Thr Thr Asn Val Asn Asn			
170	175	180	
gcg acc acc act tgc ttt gaa ggc ttc tcc aaa cgt gtc tgg aag aca			1170
Ala Thr Thr Thr Cys Phe Glu Gly Phe Ser Lys Arg Val Trp Lys Thr			
185	190	195	
tac ctg tcc aag atc act ata ttc att gaa gtt gtt gga ttc atc att			1218
Tyr Leu Ser Lys Ile Thr Ile Phe Ile Glu Val Val Gly Phe Ile Ile			
200	205	210	215
cct ctg ata ttg aat gtt tct tgt tct tct gtg gtg ctt aga acc ctc			1266
Pro Leu Ile Leu Asn Val Ser Cys Ser Ser Val Val Leu Arg Thr Leu			
220	225	230	
cgc aag cct gca aca ttg tct cag att ggg acc aat aag aaa aaa gtg			1314
Arg Lys Pro Ala Thr Leu Ser Gln Ile Gly Thr Asn Lys Lys Val			
235	240	245	
ttg aag atg atc aca gtg cat atg gca gtg ttt gtg gta tgc ttt gta			1362
Leu Lys Met Ile Thr Val His Met Ala Val Phe Val Val Cys Phe Val			
250	255	260	

cca tac aac tcc gtt ctc ttt tta tat gcc ttg gta cgc tcc caa gcc		1410
Pro Tyr Asn Ser Val Leu Phe Leu Tyr Ala Leu Val Arg Ser Gln Ala		
265	270	275
att act aat tgc tta ttg gaa agg ttt gca aag atc atg tac cca att		1458
Ile Thr Asn Cys Leu Leu Glu Arg Phe Ala Lys Ile Met Tyr Pro Ile		
280	285	290
acc ttg tgc ctt gca act ctg aat tgt tgc ttt gat cct ttt atc tat		1506
Thr Leu Cys Leu Ala Thr Leu Asn Cys Cys Phe Asp Pro Phe Ile Tyr		
300	305	310
tac ttc act ctt gaa tcc ttt cag aag tcc ttt tat atc aat aca cat		1554
Tyr Phe Thr Leu Glu Ser Phe Gln Lys Ser Phe Tyr Ile Asn Thr His		
315	320	325
ata agg atg gag tcg ctg ttt aag act gag aca cct ctg acc ccc aaa		1602
Ile Arg Met Glu Ser Leu Phe Lys Thr Glu Thr Pro Leu Thr Pro Lys		
330	335	340
cct tcc ctt cca gct atc caa gag gaa gtt agt gat caa aca aca aat		1650
Pro Ser Leu Pro Ala Ile Gln Glu Glu Val Ser Asp Gln Thr Thr Asn		
345	350	355
aat ggt ggt gaa tta atg ctg gaa tcc acc ttc taggtaccag aattgtcttt		1703
Asn Gly Gly Glu Leu Met Leu Glu Ser Thr Phe		
360	365	370

caggttcagc tacagtgtct ctatgattt tttccatg ctataaatag gagaaacaaa 1763

ttgaagctaa tgatactgag aatagagtaa tgtaccaa atgcagtc agat acatttgttt 1823

gaacactatt gtacatattc tgtttggc agtaattata ggtcaagtct aattacaaca 1883

acaaaaacag atcagccctct tctgttgagt tgactttca ttacctaaat gaccagtgg 1943

cttgactttt agtgatgtga gggttat taaacttaaa aaaaaaggca ttccagtaat 2003

tttgtaatt gggttggcc tataaatata gaacaaattc agggattatt taaaaacatc 2063

tgtttacta ctgatatatg ctagtatttt ttcctttt tgaattaata ttgaatttat 2123

ttaaaaaaaaaa gaactat ttt taccta atct taataagaca tactgagaaa gagaaatgtg 2183

ttgaatttta aaatattggc aaattttacc tagattttaa aaacctaaat gaagtgttg 2243

aatgaatatg ggtggaaat ttggaattt gacaacattt acgcatttat aataaccaca 2303

attagtgtca gctttaaaaa ctttctttt aaaataattc tagaattttc atatgaaatt 2363

gttaatcctg aaagggtgcta ctatgtgcc tggcaggtat aaaatggaaa actcataaaa 2423

ttaacagtgt caat taaa aaaaaaaaaac tttaagcaac actatattat ttcttaagat 2483

tttcatttat ctttatggg ggtggggatt ggctgtaga aaatattat tcttcatttt 2543

aaatgttggg gacacattac agccagagag ctacagtatt tggcccagg tcaggagtaa 2603

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<211> 370

<212> PRT

<213> Homo sapiens

<400> 174

Met Gly Asp Arg Arg Phe Ile Asp Phe Gln Phe Gln Asp Leu Asn Ser

1

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10

15

Ser Leu Arg Pro Arg Leu Gly Asn Ala Thr Ala Asn Asn Thr Cys Ile

20

25

30

Val Asp Asp Ser Phe Lys Tyr Asn Leu Asn Gly Ala Val Tyr Ser Val

35

40

45

Val Phe Ile Leu Gly Leu Ile Thr Asn Ser Ala Ser Leu Phe Val Phe

50

55

60

Cys Phe Arg Met Lys Met Arg Ser Glu Thr Ala Thr Phe Ile Thr Asn

65

70

75

80

Leu Ala Leu Ser Asp Leu Leu Phe Val Cys Thr Leu Pro Phe Lys Ile

85

90

95

Phe Tyr Asn Phe Asn Arg His Trp Pro Phe Gly Asp Thr Leu Cys Lys

100

105

110

Ile Ser Gly Thr Ala Phe Leu Thr Asn Ile Tyr Gly Ser Met Leu Phe

115

120

125

Leu Thr Cys Ile Ser Val Asp Arg Phe Leu Ala Ile Val Tyr Pro Phe

130

135

140

Arg Ser Arg Thr Ile Arg Thr Arg Arg Asn Ser Ala Ile Val Cys Ala

145

150

155

160

Gly Val Trp Ile Leu Val Leu Ser Gly Gly Ile Ser Ala Ser Leu Phe

165

170

175

Ser Thr Thr Asn Val Asn Asn Ala Thr Thr Cys Phe Glu Gly Phe

180

185

190

Ser Lys Arg Val Trp Lys Thr Tyr Leu Ser Lys Ile Thr Ile Phe Ile

195

200

205

Glu Val Val Gly Phe Ile Ile Pro Leu Ile Leu Asn Val Ser Cys Ser

210

215

220

Ser Val Val Leu Arg Thr Leu Arg Lys Pro Ala Thr Leu Ser Gln Ile
 225 230 235 240

 Gly Thr Asn Lys Lys Val Leu Lys Met Ile Thr Val His Met Ala
 245 250 255

 Val Phe Val Val Cys Phe Val Pro Tyr Asn Ser Val Leu Phe Leu Tyr
 260 265 270

 Ala Leu Val Arg Ser Gln Ala Ile Thr Asn Cys Leu Leu Glu Arg Phe
 275 280 285

 Ala Lys Ile Met Tyr Pro Ile Thr Leu Cys Leu Ala Thr Leu Asn Cys
 290 295 300

 Cys Phe Asp Pro Phe Ile Tyr Tyr Phe Thr Leu Glu Ser Phe Gln Lys
 305 310 315 320

 Ser Phe Tyr Ile Asn Thr His Ile Arg Met Glu Ser Leu Phe Lys Thr
 325 330 335

 Glu Thr Pro Leu Thr Pro Lys Pro Ser Leu Pro Ala Ile Gln Glu Glu
 340 345 350

 Val Ser Asp Gln Thr Thr Asn Asn Gly Gly Glu Leu Met Leu Glu Ser
 355 360 365

 Thr Phe
 370

<210> 175

<211> 2299

<212> DNA

<213> Homo sapiens

<220>

<221> CDS

<222> (67)..(1176)

<400> 175

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aagtcc atg ggt gac aga aga ttc att gac ttc caa ttc caa gat tca 108

Met Gly Asp Arg Arg Phe Ile Asp Phe Gln Phe Gln Asp Ser

1 5 10

aat tca agc ctc aga ccc agg ttg ggc aat gct act gcc aat aat act 156

Asn Ser Ser Leu Arg Pro Arg Leu Gly Asn Ala Thr Ala Asn Asn Thr

15 20 25 30

tgc att gtt gat gat tcc ttc aag tat aat ctc aat ggt gct gtc tac 204

Cys Ile Val Asp Asp Ser Phe Lys Tyr Asn Leu Asn Gly Ala Val Tyr

35 40 45

agt gtt gta ttc atc ttg ggt ctg ata acc aac agt gtc tct ctg ttt 252

Ser Val Val Phe Ile Leu Gly Leu Ile Thr Asn Ser Val Ser Leu Phe

50	55	60	
gtc ttc tgt ttc cgc atg aaa atg aga agt gag act gct att ttt atc 300			
Val Phe Cys Phe Arg Met Lys Met Arg Ser Glu Thr Ala Ile Phe Ile			
65	70	75	
acc aat cta gct gtc tct gat ttg ctt ttt gtc tgt aca cta cct ttt 348			
Thr Asn Leu Ala Val Ser Asp Leu Leu Phe Val Cys Thr Leu Pro Phe			
80	85	90	
aaa ata ttt tac aac ttc aac cgc cac tgg cct ttt ggt gac acc ctc 396			
Lys Ile Phe Tyr Asn Phe Asn Arg His Trp Pro Phe Gly Asp Thr Leu			
95	100	105	110
tgc aag atc tct gga act gca ttc ctt acc aac atc tat ggg agc atg 444			
Cys Lys Ile Ser Gly Thr Ala Phe Leu Thr Asn Ile Tyr Gly Ser Met			
115	120	125	
ctc ttt ctc acc tgt att agt gtg gat cgt ttc ctg gcc att gtc tat 492			
Leu Phe Leu Thr Cys Ile Ser Val Asp Arg Phe Leu Ala Ile Val Tyr			
130	135	140	
cct ttt cga tct cgt act att agg act agg agg aat tct gcc att gtg 540			
Pro Phe Arg Ser Arg Thr Ile Arg Thr Arg Arg Asn Ser Ala Ile Val			
145	150	155	
tgt gct ggt gtc tgg atc cta gtc ctc agt ggc ggt att tca gcc tct 588			
Cys Ala Gly Val Trp Ile Leu Val Leu Ser Gly Gly Ile Ser Ala Ser			
160	165	170	

ttg ttt tcc acc act aat gtc aac aat gca acc acc acc tgc ttt gaa	636		
Leu Phe Ser Thr Thr Asn Val Asn Asn Ala Thr Thr Thr Cys Phe Glu			
175	180	185	190
ggc ttc tcc aaa cgt gtc tgg aag act tat tta tcc aag atc aca ata	684		
Gly Phe Ser Lys Arg Val Trp Lys Thr Tyr Leu Ser Lys Ile Thr Ile			
195	200	205	
ttt att gaa gtt gtt ggg ttt atc att cct cta ata ttg aat gtc tct	732		
Phe Ile Glu Val Val Gly Phe Ile Ile Pro Leu Ile Leu Asn Val Ser			
210	215	220	
tgc tct tct gtg gtg ctg aga act ctt cgc aag cct gct act ctg tct	780		
Cys Ser Ser Val Val Leu Arg Thr Leu Arg Lys Pro Ala Thr Leu Ser			
225	230	235	
caa att ggg acc aat aag aaa aaa gta ctg aaa atg atc aca gta cat	828		
Gln Ile Gly Thr Asn Lys Lys Val Leu Lys Met Ile Thr Val His			
240	245	250	
atg gca gtc ttt gtg gta tgc ttt gta ccc tac aac tct gtc ctc ttc	876		
Met Ala Val Phe Val Val Cys Phe Val Pro Tyr Asn Ser Val Leu Phe			
255	260	265	270
ttg tat gcc ctg gtg cgc tcc caa gct att act aat tgc ttt ttg gaa	924		
Leu Tyr Ala Leu Val Arg Ser Gln Ala Ile Thr Asn Cys Phe Leu Glu			
275	280	285	

aga ttt gca aag atc atg tac cca atc acc ttg tgc ctt gca act ctg 972
 Arg Phe Ala Lys Ile Met Tyr Pro Ile Thr Leu Cys Leu Ala Thr Leu
 290 295 300

aac tgt tgt ttt gac cct ttc atc tat tac ttc acc ctt gaa tcc ttt 1020
 Asn Cys Cys Phe Asp Pro Phe Ile Tyr Tyr Phe Thr Leu Glu Ser Phe
 305 310 315

cag aag tcc ttc tac atc aat gcc cac atc aga atg gag tcc ctg ttt 1068
 Gln Lys Ser Phe Tyr Ile Asn Ala His Ile Arg Met Glu Ser Leu Phe
 320 325 330

aag act gaa aca cct ttg acc aca aag cct tcc ctt cca gct att caa 1116
 Lys Thr Glu Thr Pro Leu Thr Thr Lys Pro Ser Leu Pro Ala Ile Gln
 335 340 345 350

gag gaa gtg agt gat caa aca aca aat aat ggt ggt gaa tta atg cta 1164
 Glu Glu Val Ser Asp Gln Thr Thr Asn Asn Gly Gly Glu Leu Met Leu
 355 360 365

gaa tcc acc ttt taggtatgag aaatgtgttc aggtccagat atggtttctc 1216
 Glu Ser Thr Phe
 370

ctataatttt tcctatgcta taaaactaaag atttgaagct aatgatactg agaataatgc 1276

accaaatcca gtcagataca ttgtttgaa ggtatactgt agagtttta ttgctgttt 1336

gttcagtaat tataggtaaa atctaattac aacaaccaag atggattgcc aaactcttct 1396

gcttggttgg aatttcattt tatgcattt tccaggtggc tagtggcatt tgataatata 1456

gagatgactt tgaaactttc aaaaaggat ttctattcca atgatattt gtaatttagt 1516

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<213> Homo sapiens

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Ser Leu Arg Pro Arg Leu Gly Asn Ala Thr Ala Asn Asn Thr Cys Ile

20

25

30

Val Asp Asp Ser Phe Lys Tyr Asn Leu Asn Gly Ala Val Tyr Ser Val

35

40

45

Val Phe Ile Leu Gly Leu Ile Thr Asn Ser Val Ser Leu Phe Val Phe

50

55

60

Cys Phe Arg Met Lys Met Arg Ser Glu Thr Ala Ile Phe Ile Thr Asn

65

70

75

80

Leu Ala Val Ser Asp Leu Leu Phe Val Cys Thr Leu Pro Phe Lys Ile

85

90

95

Phe Tyr Asn Phe Asn Arg His Trp Pro Phe Gly Asp Thr Leu Cys Lys

100	105	110
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Ile Ser Gly Thr Ala Phe Leu Thr Asn Ile Tyr Gly Ser Met Leu Phe

115	120	125
-----	-----	-----

Leu Thr Cys Ile Ser Val Asp Arg Phe Leu Ala Ile Val Tyr Pro Phe

130	135	140
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Arg Ser Arg Thr Ile Arg Thr Arg Arg Asn Ser Ala Ile Val Cys Ala

145	150	155	160
-----	-----	-----	-----

Gly Val Trp Ile Leu Val Leu Ser Gly Gly Ile Ser Ala Ser Leu Phe

165	170	175
-----	-----	-----

Ser Thr Thr Asn Val Asn Asn Ala Thr Thr Thr Cys Phe Glu Gly Phe

180	185	190
-----	-----	-----

Ser Lys Arg Val Trp Lys Thr Tyr Leu Ser Lys Ile Thr Ile Phe Ile

195	200	205
-----	-----	-----

Glu Val Val Gly Phe Ile Ile Pro Leu Ile Leu Asn Val Ser Cys Ser

210	215	220
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Ser Val Val Leu Arg Thr Leu Arg Lys Pro Ala Thr Leu Ser Gln Ile

225	230	235	240
-----	-----	-----	-----

Gly Thr Asn Lys Lys Lys Val Leu Lys Met Ile Thr Val His Met Ala

245	250	255
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Val Phe Val Val Cys Phe Val Pro Tyr Asn Ser Val Leu Phe Leu Tyr

260

265

270

Ala Leu Val Arg Ser Gln Ala Ile Thr Asn Cys Phe Leu Glu Arg Phe

275

280

285

Ala Lys Ile Met Tyr Pro Ile Thr Leu Cys Leu Ala Thr Leu Asn Cys

290

295

300

Cys Phe Asp Pro Phe Ile Tyr Tyr Phe Thr Leu Glu Ser Phe Gln Lys

305

310

315

320

Ser Phe Tyr Ile Asn Ala His Ile Arg Met Glu Ser Leu Phe Lys Thr

325

330

335

Glu Thr Pro Leu Thr Thr Lys Pro Ser Leu Pro Ala Ile Gln Glu Glu

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Thr Phe

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Leu Leu Arg Leu Leu Val Leu Gly Leu Trp Leu Ala Leu Leu Arg Ser

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Val Ala Gly Glu Gln Ala Pro Gly Thr Ala Pro Cys Ser Arg Gly Ser

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Ser Trp Ser Ala Asp Leu Asp Lys Cys Met Asp Cys Ala Ser Cys Arg

45 50 55

gcg cga ccg cac agc gac ttc tgc ctg ggc tgc gct gca gca cct cct 245

Ala Arg Pro His Ser Asp Phe Cys Leu Gly Cys Ala Ala Ala Pro Pro

60 65 70

gcc ccc ttc cgg ctg ctt tgg ccc atc ctt ggg ggc gct ctg agc ctg 293

Ala Pro Phe Arg Leu Leu Trp Pro Ile Leu Gly Gly Ala Leu Ser Leu

75

80

85

acc ttc gtg ctg ggg ctg ctt tct ggc ttt ttg gtc tgg aga cga tgc 341

Thr Phe Val Leu Gly Leu Leu Ser Gly Phe Leu Val Trp Arg Arg Cys

90

95

100

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Arg Arg Arg Glu Lys Phe Thr Thr Pro Ile Glu Glu Thr Gly Gly Glu

105

110

115

120

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125

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Leu Trp Leu Ala Leu Leu Arg Ser Val Ala Gly Glu Gln Ala Pro Gly

20

25

30

Thr Ala Pro Cys Ser Arg Gly Ser Ser Trp Ser Ala Asp Leu Asp Lys

35

40

45

Cys Met Asp Cys Ala Ser Cys Arg Ala Arg Pro His Ser Asp Phe Cys

50

55

60

Leu Gly Cys Ala Ala Pro Pro Ala Pro Phe Arg Leu Leu Trp Pro

65

70

75

80

Ile Leu Gly Gly Ala Leu Ser Leu Thr Phe Val Leu Gly Leu Leu Ser

85

90

95

Gly Phe Leu Val Trp Arg Arg Cys Arg Arg Glu Lys Phe Thr Thr

100

105

110

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115

120

125

Gln

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【0129】

【配列表フリーテキスト】

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配列番号180：プライマー

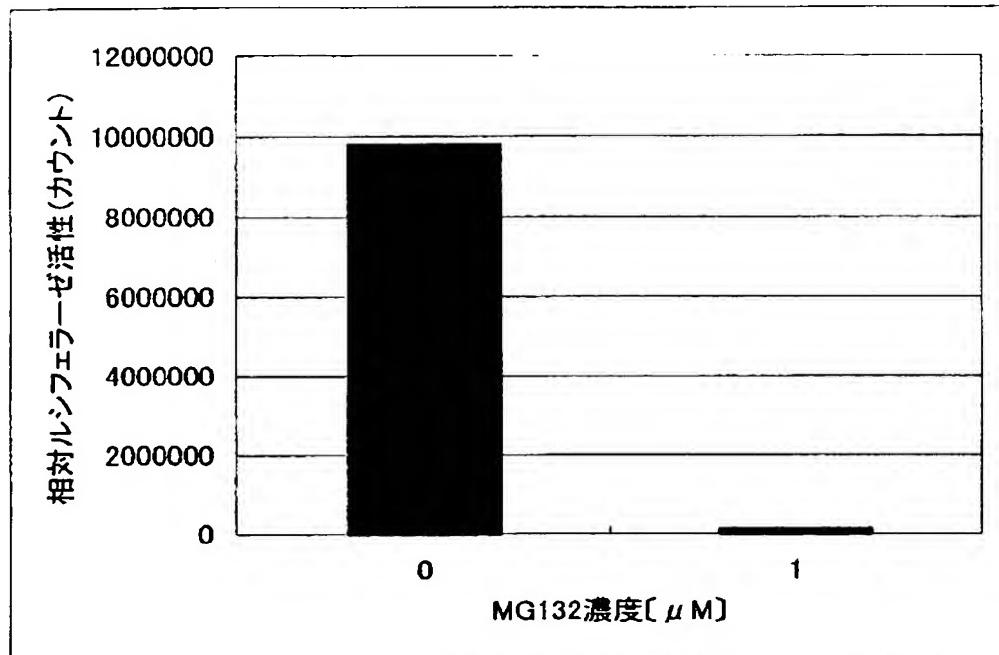
【図面の簡単な説明】

【図1】

図1は、実施例3のプロテアソーム阻害剤MG132によるNF- κ Bのレポーター活性抑制を示す図である。図中で横軸は、MG132濃度、縦軸は、相対ルシフェラーゼ活性を示す。

【書類名】 図面

【図1】



【書類名】 要約書

【要約】

【課題】 N F - κ B の過剰な活性化または阻害が関与する疾患の診断、治療または予防等に使用される N F - κ B 作用を有するタンパク質の提供。

【解決手段】 ヒト肺線維芽細胞から作製した c D N A ライブラリーから、プラスミド p N F κ B - L u c を用いて、 N F - κ B を活性化する作用を有するタンパク質をコードする c D N A をクローニングして、その D N A 配列およびそれより推定されるアミノ酸配列を決定した。同タンパク質、これをコードする D N A 、同 D N A を含有する組換えベクターおよび同組換えベクターを含有する形質転換体は、 N F - κ B の活性化を阻害または促進する物質のスクリーニングに使用される。

【選択図】 なし

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【提出日】 平成13年11月12日

【あて先】 特許庁長官 殿

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【出願番号】 特願2001-254018

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【手続補正 1】

【補正対象書類名】 特許願

【補正対象項目名】 発明者

【補正方法】 変更

【補正の内容】

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1. 変更年月日 2001年 1月 4日

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